

Qy 388 ATATGGGCACCTAGCTGCATGAATAATATGATGGAAGAATGGATCAAGTGGAGTTTCCATA 447

Dh 241 ATATGGGCGCTGGCTGCATGAATAAGGACGGAAGAATGGATCAGTGGAGTTTCCATA 300

| | | | |
|----|------|--|------|
| Qy | 448 | GCTATGAAACTTATCAAACTGAAGCTACAAGGATATCAGCTACCCCTCTGCACCTTCCCCCT | 507 |
| Db | 301 | GCCATGAAGCTCATCAAACTGAAGCTCAAGGATATCAGCTCCCCCGCACCTTCCGCCCT | 360 |
| Qy | 508 | GTCAATGAACAGCAACCAAGTTGCTATTCTTCTAGCCGACAGGATTTGATATGGAGGTATC | 567 |
| Db | 361 | GTCAATGAAGCAGCAGCCAGCGGCCATCTCTAGTGCACCAAGCGTTTGGTATAGGAGGGATG | 420 |
| Qy | 568 | GCCAGCATGCCACCGCTTACAGCTGTGTCTCCAGTGCCAATGGGATCCATTTCAGATTGTT | 627 |
| Db | 421 | GCTGGATGCCACCACTGCACAGCTTGTCTCCGTGCAATGGCTCCATCCCACTGTTT | 480 |
| Qy | 628 | GGAAATGTTCCAAACCCCTAGTATCTTCTGTTCCTCCACAGAGCTGTGCCCCCTGGCTTAAC | 687 |
| Db | 481 | GGAAATGTTCCGCCCTTAGTATCTTCTGTCTCCCTCAAGCAGCAGTGCTCCCTGGCTTAAC | 540 |
| Qy | 688 | GGGGCTCCCCCTGTTATACAACCTCTGCCTGCATTTGCTCATCTCGCAGCCACATTTGCCA | 747 |
| Db | 541 | GGGGCCCCCTCTGTGCATACAGCCTCTGCCTGCATTTGCTCATCTCGCAGCCACATTTGCCA | 600 |
| Qy | 748 | AAGAGTTCTTCTCTTTAGTAGATCTGGTCCAGGGTCACAACTAAACACTAAATTACAAAAG | 807 |
| Db | 601 | AAGAGTTCTTCTTTCACAGATCTGGTCCAGGGTCACAACTAAACACTAAATTACAGAAG | 660 |
| Qy | 808 | GCA CAGTCAATTTGATGTGGCCAGTGTCCCAACAGTGGCAGAGTGGGCTGTTCCTCAGTCA | 867 |
| Db | 661 | GCA CAAATCATTTGATGTAGCCAGCGCCCCCTGCAGCGCAGAAATGGGCTGTGCTCTCAGTCG | 720 |
| Qy | 868 | TCAGAGCTGAAATCAGGCAATTAATCAATAGTCTATGACAACTATGAGTGAGTGACACTTA | 927 |
| Db | 721 | TCAGAGCTGAAATCAGGCAATTAATCAACAGTCCAGCAAGACCATGAGTGAGCACTTA | 780 |
| Qy | 928 | ACAGGTCCCCAAGCAAGAACTATTCTTATG CAGTCAAGTTTACCAAGGCTCAGCTGGCT | 987 |
| Db | 781 | ACAGGTCCCCAGGCAAGAACTATTCTCATGCAATCAAGTTTACCCAGGCTCAGCTGGCT | 840 |
| Qy | 988 | TCAATATGGAATCTTTCTGCATTTGATCAAGATGGGAAAACTTACAGCAGAGGAATTTATC | 1047 |
| Db | 841 | TCAATCTGGAATCTTTCCGACATTTGATCAAGATGGAAGGCTCACCGCAGGAAGAAATTCA | 900 |
| Qy | 1048 | CTGCAATGCACTTCATTGTATGTAGTATGTCTGGCCAAACACATGCCACCTGTCTGCGCT | 1107 |
| Db | 901 | CTAGCGATGCACTGATTTGATGTCTGATGTCTGCTCAGCCATCTGCGCTGTGCTGCTCT | 960 |
| Qy | 1108 | CCAGAAATACATTCACCTCTCTTTTAGAAGAGTTCCGATCTGGCAGTGGTATATCTGTCAAT | 1167 |
| Db | 961 | CCAGAAATACATCTCTCTCTCTTTTAGAAGAGTTCCGCTCTGGCAGTGGGATGTCGCTCAT | 1020 |
| Qy | 1168 | AGCTCAACATCTGTAGATTCAGAGGCTTACAGAGGAACAGTTTTTAGAAGATGAACAACAA | 1227 |
| Db | 1021 | AGCTCTTCTGCTGTGAGACCAAGCGGCTGCCGAGAGGCCATCGTCAGAGGATGAGCAGCAG | 1080 |
| Qy | 1228 | CAATTAGAAAAAATAATACCTGTAACTGTTTGAAGATAAGAAAGCGGAGAACTTTTGAACGT | 1287 |
| Db | 1081 | G---TGAAAAAAGAGCTGCTGTGTGACATTTTGAAGATTAAGAAAGCGGAGAACTTCGACGA | 1137 |
| Qy | 1288 | GGCAACCTGGAACTGGAGAAACGAAGGCAAGCTCTCTCTGGAAACAGCAGCGCAAGGACAG | 1347 |
| Db | 1138 | GGCAACCTGGAGCTGGAGAACGCGGCTGCCGAGAGGCCATCGTCAGAGGATGAGCAGCAG | 1197 |
| Qy | 1348 | GAGCGCTGGCCAGCTGGAGCGCGGCGGAGCAGAGAGGAGGAGCTGTGAGCCGAGGAG | 1407 |
| Db | 1198 | GAGCGGCTGGCTAGCTGTGAGCGCGCGAGCAGAGAGGAAAGAGCGGAGCGCCAGGAA | 1257 |
| Qy | 1408 | CAAGCGCGCAAAAGACAACTGGAACCTGGAGAAGCAACTGGAAGAAAGCAGCGGAGCTAGAA | 1467 |
| Db | 1258 | CAGAGCGCAGAGACAGCTAGAGCTGGAGNAGCAGCTAGNAAAGCAGCGGAGCTGGAG | 1317 |
| Qy | 1468 | CGGCAGAGAGAGGAGGAGGAGAAAGAAATTCAGAGGCGGAGAGGCTGCAAAACCGGAA | 1527 |
| Db | 1318 | AGGCAGAGAGAGGAGGAGAAAGAGAGAGATTCAGAGGCGGAGGAGGAGCAAAACCGGAG | 1377 |

| | | | |
|----|------|---|------|
| QY | 1528 | CTTGAAAGGCAACGACAACTTTGAGTGGGAACGGAATCGAAGGCCAAGNACTACTTAATCAA | 158 |
| DB | 1378 | CTCGAACGGCAGGCACAGCTTTGAATGGGAACGGAACCGGAGACAGGAACTCCTGACTCAG | 1437 |
| QY | 1588 | AGAAACAAAGAACAAAGAGGACATAGTTGTACTGAAAGCAAAAGAAAAAGACTTTGGGAATTT | 1647 |
| DB | 1438 | AGGAACAAGACACAGGAGGCATCTGTGGTTCTTGAGGCGAGGAGGAAGACTCTTGGAGTTT | 1497 |
| QY | 1648 | GAATTAGAAGCTCTTAATTGATAAAAAAGCATCAACTAGNAGGGAAACTTCAAGATATCAGA | 1707 |
| DB | 1498 | GAATTAGAAGCTCTGAATGACAAAAGACATCAGCTGGAAGGAAAACTTCAGGATATCAGG | 1557 |
| QY | 1708 | TGTCGATTGACCAACCCAAAGGCAAGAAATTTGAGAGACACAAACAAATCTAGAGAGTTGAGA | 1767 |
| DB | 1558 | TGTCGACTGGCAACCCCAAGAGCCAAAGAAATTTGAGAGACAAACAAAGCTAGAGAGCTGAGA | 1617 |
| QY | 1768 | ATTGCCGAATACCCCATCTTACAGCAACAAATTAACAGNAATCTCAGCAAAATGCTTGGGAAGA | 1827 |
| DB | 1618 | ATTGCGGAATACCCCATCTTACAGCAACAGTTGCAGGAATCTTCAGCAGATGCTTTGGHAGA | 1677 |
| QY | 1828 | CTTATTCCAGAAAAACAGATACTCAATGACCAATTTAAAAACAAGTTTCAGCAGAAACAGTTTG | 1887 |
| DB | 1678 | CTTATTCCAGAAAGCAGATACTCTAGTGACCGATTAAACACAAGTCCAGCAGACACAGCTTG | 1737 |
| QY | 1888 | CACAGAGATTCACTTTGTTCACCTTAAAGAGCCTTTAGAAGCAAAAGAAATCTAGCTCGGCAG | 1947 |
| DB | 1738 | CATAGAGATTTCGCTTCTTACCCCTCAAAGAGCCTTTGGAAGCAAAAGGAACTGGGCCCGCAG | 1797 |
| QY | 1948 | CACCTACGAGCCAACTGGGATGAAGTGGGAGAAAGNAACTAGATCNAAACTACACAGGAGATT | 2007 |
| DB | 1798 | CAGCTTCGAGAGCAGCTTGGACGAGGTTGGAAAGAGACCAGGTTCAAAGCTGCGAGGAGATT | 1857 |
| QY | 2008 | GATATTTTCAATTAATCAGCTGAAGGAACTAAGAGAAATACACAAATAGCAACAACTCCAG | 2067 |
| DB | 1858 | GATGTTTTCAACACAGCTTGAAGAACTCAGAGAGNATACACAGTAAACAGCAGCTCCAG | 1917 |
| QY | 2068 | AAGCAAAAGTCCATGGAGGCTGAACGACTGAATCAAGAAAGAAACAAGAAAGAAATCATTA | 2127 |
| DB | 1918 | AAGCAGAGGTTCCATTCGAGGCGGAGAGGCTTAAGCAGAAAGAGCAGGAGAGGAAGAGCCTG | 1977 |
| QY | 2128 | GAATTAGAAAAACAAAAGAGAAAGCCAAAGACGACTCAGGAAAGGGACACAGCAGTGG | 2187 |
| DB | 1978 | GAGTTGGAGAAAGCAAAAGSAGAAAGGTCAGAGACGAGTTTCAGGAAAGGGACACAGCAATGG | 2037 |
| QY | 2188 | CTGAGCATGTGCAGCAGGAGGACGAGCATCAGAGACCAAGAAAACTCCACAGAGAGGAA | 2247 |
| DB | 2038 | CAGAGCATGTCCAGAGAG--GAGCAGCAGCGCCCTCGHAAACCCACAGGAGGAC | 2094 |
| QY | 2248 | AAACTGAAAAGGGAGGAGTGTCGAGAGAGGATGTCGAGAGAGGATGTCGAGAGAGGAAATGTA | 2307 |
| DB | 2095 | AAACTGAAAAGGGAAAGACAGTGTCAAGAAAGAGGAGGCGGAAGAGAGAGCCAAAGCCGAA | 2154 |
| QY | 2308 | GCAACAGACAAGCTGGGTCCGCTTTTCCATCAACACCAAGAAACGAGTAAAGCCAGCTGTC | 2367 |
| DB | 2155 | GTGCAAGACACAGCAGAGTCGGCTTTTCCATCCATCAACAGAGCCAGCTAAGCCGCGC--G | 2211 |
| QY | 2368 | CAGSCACCTTGCTCCACTGCAGAAAAAGGTCCACTTACCAATTTCTGCACAGGAAAAATGTA | 2427 |
| DB | 2212 | CAGGCAACCTTGGGCCCAACGACAGAAAGGTCCGCTTCAATCTCTGCACAGGAGAGTGCC | 2271 |
| QY | 2428 | AAAGTGTGTATTACCCGGCAGCTGTATCCCTTTTGAATCCAGAGACCCATGATGAATCACT | 2487 |
| DB | 2272 | AAAGTGTGTATTACCCGAGCGCTGTACCCCTTTTGGTTCAGAAAGTCATGACGAGATCACC | 2331 |
| QY | 2488 | ATCCAGCCAGAGACATAGTTCAT-----GGTGATGAAAGCCAAACTGGA | 2532 |
| DB | 2332 | ATCCAGCCAGGAGATATAGTCATGGTTTAAAGGGGGAATGGGTGGATGAAAGCCAGACCGGA | 2391 |
| QY | 2533 | GAACCCGGCTGGCTTGGAGGAGAAATTAAGGAAAGACAGGGTGGTTTCCCTGCAAACTAT | 2592 |
| DB | 2392 | GAGCCAGATGGCTTTGGAGGAGAACCGAAGGGGAAAGACAGATGGTTTCCCTGCAAACTAT | 2451 |
| QY | 2593 | GCAGAGAAAAATCCCAAGAAAAATGAGGTTTCCCGCTCCAGTGAAACCAAGTGACTGATTCAACA | 2652 |

Db 130 GAGAGTCAGTGGGAGGTTGGGCTCTCCCGTCCAAATTATGTGAAGCTGACACAGACA 189
Qy 3623 TGGACCCAAAGCCAGCAATGAATCATATGTTGTCCATCCCGCTCAGGCTTGAAAGTCTCT 3682
Db 190 TGGACCCAGCCAGCAATGAATCATATGTTGTCCAT-CCCCCTCAGGCTTGAAAGCTCT 248
Qy 3683 CAAAGAGACCCACTATCCCATATCACTGCCAGAGAGGATGATGGGAGATGCGACCTTGAT 3742
Db 249 CAAAGAGACCCACTATCCCATATCACTGCCAGAGAGGATGATGGGAGATGCGACCTTGAT 308
Qy 3743 CATGTGACTCCAGCATGATCACTACTGCTCTCTGAGTAGAGAACTCACTGCGAGCA 3802
Db 309 CATGTGACTCCAGCATGATCACTACTGCTCTCTGAGTAGAGAACTCACTGCGAGCA 368
Qy 3803 GTTTACTCTATTTTACCTTAGTTGTCATGTGATCGCAATGTTTGAGTTATTACTTGCAGAG 3862
Db 369 GTTTACTCTATTTTACCTTAGTTGTCATGTGATCGCAATGTTTGAGTTATTACTTGCAGAG 427
Qy 3863 ATAGGAGCAAAATTAACAAAACACACAGGGTAGTGGGCTCTTTTGTGGCTTTCTTAGTT 3922
Db 428 GCGAAGAC--AAATTGCAAGAACTGCACAGGGTGGTGGTCTTTTGT-TCCTTCTTAGTC 484
Qy 3923 ACTCAATTCAGCTTTCCCGCCACTTTGACAGGTGCTTTCAATAGTTTAAATATTTT 3982
Db 485 ACTCAGACTGACGGGCCCCGCTTCASWGWAGYGACTWKSAA-AGYVYAARAKKAYTTT 543
Qy 3983 TAAATATATATTTAGCTTTTAAATAAACA 4012
Db 544 TAAATGTGATTTTACGCTTTTAAATAAACA 573

RESULT 3
US-11-091-883-433
; Sequence 433, Application US/11091883
; Publication No. US20060024693A1
; GENERAL INFORMATION:
; APPLICANT: CIBELLI, JOSE
; APPLICANT: FERNANDEZ, EMILIO O.
; APPLICANT: JORDAO DE MEGALHAES, GUILHERME
; APPLICANT: KOCABAS, ARIF
; APPLICANT: CROSBY, JAVIER A.
; TITLE OF INVENTION: IDENTIFICATION OF GENES OR POLYPEPTIDES THE EXPRESSION OF WHICH
; TITLE OF INVENTION: CORRELATES TO FERTILITY, OVARIAN FUNCTION AND/OR FETAL/NEWBORN
; FILE REFERENCE: 53942US
; CURRENT APPLICATION NUMBER: US/11/091,883
; PRIOR FILING DATE: 2005-03-29
; PRIOR APPLICATION NUMBER: 60/556,875
; PRIOR FILING DATE: 2004-03-29
; NUMBER OF SEQ ID NOS: 513
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 433
; LENGTH: 4053
; TYPE: DNA
; ORGANISM: Homo sapiens
US-11-091-883-433

Query Match 3.2%; Score 165.2; DB 11; Length 4053;
Best Local Similarity 60.4%; Pred. No. 1.1e-25;
Matches 343; Conservative 0; Mismatches 193; Indels 32; Gaps 3;
Qy 3208 GAAATTTCCAGGTTATTCCTCATACCGCCACCGGCCCCGAGCAGTCTCACTCTCGCC 3267
Db 334 GAGATTGCTCAGGTAACCTTCAGCATATGTTGCTTCTGCTTCTGAACTTCACTGCTTGA 393
Qy 3268 CCTGGTCAGCTGATTTTGTATCCGAAAGAACCCAGGTGGATGTTGGGAGAGAGCTG 3327
Db 394 CCAGGACAGTTATATTAATTTTAAAGAAAAATACAAAGTGGGTTGGGCAAGAGAGTTA 453
Qy 3328 CAAGCACGTGGGAAAAAGCGCCAGATAGGCTGCTCCAGCTAATTTATGTAAAGCTTCTA 3387
Db 454 CAGGCCAGAGAAAAAGCGACAGAAAGGATGTTTCTTCCAGTCAATGTTAAACTTTTG 513

Qy 3388 AGCCCTGGGACGAGCAAAATCACTCCAAACAGAGCCACCCTAAAGTCAACAGCATTAGCGGCA 3447
Db 514 GGTCAAGCAGGTGAAAGAGCCA-----CACCTGCTTTTCATCTCT 552
Qy 3448 GTGTCCAGGTCATTTGGGATGTACGACTACACCGGCGAGAAATGACGATGAGCTGCGCTTC 3507
Db 553 GTATGTCAAGGTGATTTGCTATGTATGACTATGCGAGCAAAATATGAAGATGAGCTCAGTTTC 612
Qy 3508 AACAAAGGCCAGATCATCAACGCTCTCAACAAAGGAGGACCCCTGACTGGTGGAAAGGAGAA 3567
Db 613 TCCAAAGGACAACTCATTAATGTTATGAACAAAGATGATCTGATTTGGTGGCAAGAGAG 672
Qy 3568 GTCAATGACAAAGTGGGGCTCTTCCATCCCAATATGTGAAGCTGACACAGACATGGAC 3627
Db 673 ATCAACGGGGTACTGCTCTCTTTCTTTCAAACTACGTTAAGATGACGACAGACTCAGAT 732
Qy 3628 CCAAGCCAGCAATGAATCATATATGTTGTCATCCCGCTCAGGCTTGAAGTCTCTCAAG 3687
Db 733 CCAAGTCAACAGTGACCCA-----ATGTTGCTTCCAGTTGTGAAGCACCCAG 782
Qy 3688 AGACCCACTATCCCATATCACTGCCAGAGGAGATGATGGGAGATGACGCTTTGATCATGT 3747
Db 783 AGACCCACTAT-CCAAGTTTCACTCTAGCTGGAGGCGAGGCGAGCCCTGATCAAAAT 841
Qy 3748 GACTTCCAGCATGATCACTACTGCTT 3775
Db 842 ATCTCTACAAATTCGTTTACTTCGTT 869

RESULT 4

US-10-750-185-31648
; Sequence 31648, Application US/10750185
; Publication No. US20050260603A1
; GENERAL INFORMATION:
; APPLICANT: MMI GENOMICS, INC.
; APPLICANT: DENISE, Sue K.
; APPLICANT: KERR, Richard
; APPLICANT: ROSENFELD, David
; APPLICANT: HOLM, Tom
; APPLICANT: BATES, Stephen
; APPLICANT: FANTIN, Dennis
; TITLE OF INVENTION: COMPOSITIONS FOR INFERRING BOVINE TRAITS
; FILE REFERENCE: MM1100-2
; CURRENT APPLICATION NUMBER: US/10/750,185
; CURRENT FILING DATE: 2003-12-31
; PRIOR APPLICATION NUMBER: US 60/437,482
; PRIOR FILING DATE: 2002-12-31
; NUMBER OF SEQ ID NOS: 64922
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 31648
; LENGTH: 874
; TYPE: DNA
; ORGANISM: Bovine 19866880882217
US-10-750-185-31648

Query Match 1.9%; Score 96.2; DB 7; Length 874;
Best Local Similarity 69.3%; Pred. No. 4.2e-11;
Matches 131; Conservative 0; Mismatches 58; Indels 0; Gaps 0;
Qy 2917 CAGGCTGAAAGGTGAGGGGCTCAAGCTCAAGCCCTATATCTTTGGAGAGCCAAAAA 2976
Db 101 CAGGACAGGTTGTAGAAAACCTGAAAGCACAGGCCCTTTGTTCTTGAGCTGCAAGAAA 160
Qy 2977 GACAAACCACTTAAATTTTAAACAAAATGATGTCACTCAGCGTCTCTGGAACAGCAAGCATG 3036
Db 161 GAAACCACTTGAACCTTCTCAAAACATGATATTAATTAATGCTTAGAGCAGCAAGAAAT 220
Qy 3037 TGGTGGTTGGAGAAAGTTCAAGGTGAGGGTTGGTTCCCAAGCTTTACGTGAAACTC 3096
Db 221 TGGTGGTTGGGAGGTTTACGGAGGAGAGATGGTTTCCGAAATCTTACGTCAAAATC 280
Qy 3097 ATTTTCAGG 3105

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Db      281 ATCCCTGGG 289
|| | | |
Best Local Similarity 51.5%; Pred. No. 6.7e-06;
Matches 177; Conservative 0; Mismatches 167; Indels 0; Gaps 0;

RESULT 5
US-10-750-623-31648
; Sequence 31648, Application US/10750623
; Publication No. US20050287531A1
; GENERAL INFORMATION:
; APPLICANT: MMI GENOMICS, INC.
; APPLICANT: DENISE, Sue K.
; APPLICANT: KERR, Richard
; APPLICANT: ROSENFELD, David
; APPLICANT: HOLM, Tom
; APPLICANT: BATES, Stephen
; APPLICANT: FANTIN, Dennis
; TITLE OF INVENTION: METHODS AND SYSTEMS FOR INFERRING BOVINE TRAITS
; FILE REFERENCE: MM1100-1
; CURRENT APPLICATION NUMBER: US/10/750,623
; CURRENT FILING DATE: 2003-12-31
; PRIOR APPLICATION NUMBER: US 60/437,482
; PRIOR FILING DATE: 2002-12-31
; NUMBER OF SEQ ID NOS: 64922
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 31648
; LENGTH: 874
; TYPE: DNA
; ORGANISM: Bovine 19866880882217
US-10-750-623-31648

Query Match      1.9%; Score 96.2; DB 7; Length 874;
Best Local Similarity 69.3%; Pred. No. 4.2e-11;
Matches 131; Conservative 0; Mismatches 58; Indels 0; Gaps 0;

QY      2917 CAGGTCGAAAGGTGGAGGGCTACAGCTCAAGCCCTATATCTTGGAGAGCAAAAAA 2976
Db      101 CAGGACAGGTTGTAGAAAACCTGAAGCACAGGCCCTTTGTTCTGGACTGCAAGAAA 160

QY      2977 GACAACACATTAATATTTTAAACAAAATGATGTATCATCACCGTCTCTGGAAACAGCAGACATG 3036
Db      161 GAAAACCATTTGAACCTTCTCAAAACATGATATTATTACTGTCTTAGAGCAGCAAGAAAT 220

QY      3037 TGGTGGTTTGGAGAGTTCAAGTCACAGGGTTGGTCCCAAGTCTTACGTGAACATC 3096
Db      221 TGGTGGTTTGGGGAGGTTTCACGGAGGAAGAGGATGGTTCCGAAATCTTACGTCAAAATC 280

QY      3097 ATTTTCAGGG 3105
Db      281 ATCCCTGGG 289

RESULT 6
US-11-121-086-5
; Sequence 5, Application US/11121086
; Publication No. US20050266459A1
; GENERAL INFORMATION:
; APPLICANT: POULSEN, TIM S.
; APPLICANT: NIELSEN, KIRSTEN V.
; TITLE OF INVENTION: NUCLEIC ACID PROBES AND NUCLEIC ACID ANALOG PROBES
; FILE REFERENCE: 09138.6000-00000
; CURRENT APPLICATION NUMBER: US/11/121,086
; CURRENT FILING DATE: 2005-05-04
; PRIOR APPLICATION NUMBER: 60/567,570
; PRIOR FILING DATE: 2004-05-04
; NUMBER OF SEQ ID NOS: 107
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 5
; LENGTH: 153376
; TYPE: DNA
; ORGANISM: Homo sapiens
US-11-121-086-5

Query Match      1.5%; Score 76.8; DB 11; Length 153376;

Best Local Similarity 51.5%; Pred. No. 6.7e-06;
Matches 177; Conservative 0; Mismatches 167; Indels 0; Gaps 0;

QY      1183 GATCAGAGGCTACCAAGAACCCAGTTTGTAGAGATGAACAACAACATTTAGNAAGAAA 1242
Db      15959 GAGGAGAAAGGAGGAGGAGAAACAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 16018

QY      1243 TTACCTGTAAACGTTTGAAGATAAGAGCGGAGAGACTTTTGAACGTGGCAACCTTGGAACTG 1302
Db      16019 TAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 16078

QY      1303 GAGAAACGAAAGCAAGCTCTCTTGGAAACAGCAGCGGCAAGGAGGAGGAGGAGGAGGAGGAG 1362
Db      16079 GAGGAAACAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 16138

QY      1363 CTGGAGCGGCGGAGCAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 1422
Db      16139 GAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 16198

QY      1423 CAACTGGAACCTGGAGAAAGCAACTGGAAAGACAGCGGAGCTAGAACGGCAGAGAGGAGGAG 1482
Db      16199 GAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 16258

QY      1483 GAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 1526
Db      16259 GAACAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGA 16302

RESULT 7
US-10-750-185-31653
; Sequence 31653, Application US/10750185
; Publication No. US20050260603A1
; GENERAL INFORMATION:
; APPLICANT: MMI GENOMICS, INC.
; APPLICANT: DENISE, Sue K.
; APPLICANT: KERR, Richard
; APPLICANT: ROSENFELD, David
; APPLICANT: HOLM, Tom
; APPLICANT: BATES, Stephen
; APPLICANT: FANTIN, Dennis
; TITLE OF INVENTION: COMPOSITIONS FOR INFERRING BOVINE TRAITS
; FILE REFERENCE: MM1100-2
; CURRENT APPLICATION NUMBER: US/10/750,185
; CURRENT FILING DATE: 2003-12-31
; PRIOR APPLICATION NUMBER: US 60/437,482
; PRIOR FILING DATE: 2002-12-31
; NUMBER OF SEQ ID NOS: 64922
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 31653
; LENGTH: 2170
; TYPE: DNA
; ORGANISM: Bovine 19866880980841
US-10-750-185-31653

Query Match      1.4%; Score 74; DB 7; Length 2170;
Best Local Similarity 67.5%; Pred. No. 3.9e-06;
Matches 104; Conservative 0; Mismatches 50; Indels 0; Gaps 0;

QY      986 CTTCAATATGGAATCTTTCTGCATATGATCAAGATGGAAGAACTTACAGCAGAGGAATTTA 1045
Db      1466 CTTTCTTCAGGACTCTGGCTGACATCGATCGTGTATGGACAGCTGAAAGCTGAGGAGTTCA 1525

QY      1046 TCCTGGCAATGCACTCATTTGATGTAGCTATGTCTGCCCAACCACTGCCACCTCTCTCTGC 1105
Db      1526 TCCTTGGATGCACTCATTTGATGTAGCTATGTGCCCAAGCTGACAGCCGTTGCCGCTTCTTAC 1585

QY      1106 CTCAGAAATACATTTCCACCTTCTTTTAGAAGAGT 1139
Db      1586 CTCCTGAGCTTGTCCTCCCTCCATCTTTTAGGTGAGT 1619

RESULT 8
US-10-750-623-31653
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; CURRENT APPLICATION NUMBER: US/11/136,527
; CURRENT FILING DATE: 2005-05-25
; PRIOR APPLICATION NUMBER: US 60/574,294
; PRIOR FILING DATE: 2005-05-26
; NUMBER OF SEQ ID NOS: 362830
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 3376
; LENGTH: 2380
; TYPE: DNA
; ORGANISM: Rattus norvegicus
US-11-136-527-3376

Query Match          1.2%; Score 64.2; DB 11; Length 2380;
Best Local Similarity 55.1%; Pred. No. 0.00052;
Matches 147; Conservative 0; Mismatches 118; Indels 2; Gaps 1;

QY 2066 AGAAGCAAAAGTCCATGGAGGCTGAACGACTGAAACAGAAAGAAACAAAGAACGAAAGATCA 2125
DB 1512 AGAGGAAGAGAGGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 1453
QY 2126 TAGAATTAGAAAACAAAAGAAAGAAAGCCCAAGAGCGAGCTCAGGAAAGGCAAGCAGT 2185
DB 1452 AAGCAAGAAAGAAAGAAAGAAAGAAAGAAAGAAAG--AAGAAAGAAAGAGCAGCAGC 1395
QY 2186 GCGTGGAGCAGTGTGACGAGGAGGAGCAGCATCAGAGCAACCAAGAAACTCCACGAAGAG 2245
DB 1394 AGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAG 1335
QY 2246 AAAAAGTGAAGAGGAGGAGAGTGTCAAAAGAAAGAGTGGCGAGGAGAAAGGCAACAGG 2305
DB 1334 AGAAGAAAGAAAGAAAGAAAGAAAGAAAGAAAGAAAGAAAGAAAGAAAGAAAG 1275
QY 2306 AAGCAAGCAAGCAGTGGTGGCTTT 2332
DB 1274 AAGAGAGTCCAGCAGTGTGGTCTTT 1248

RESULT 12
US-11-136-527-155
; Sequence 155, Application US/11/136527
; Publication No. US2005028750A1
; GENERAL INFORMATION:
; APPLICANT: Wyeth
; APPLICANT: Mounts, William M
; TITLE OF INVENTION: Probe Arrays For Expression Profiling of Rat Genes
; FILE REFERENCE: 031896-041000 (AM101086)
; CURRENT APPLICATION NUMBER: US/11/136,527
; CURRENT FILING DATE: 2005-05-25
; PRIOR APPLICATION NUMBER: US 60/574,294
; PRIOR FILING DATE: 2005-05-26
; NUMBER OF SEQ ID NOS: 362830
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 155
; LENGTH: 3456
; TYPE: DNA
; ORGANISM: Rattus norvegicus
US-11-136-527-155

Query Match          1.2%; Score 63.8; DB 11; Length 3456;
Best Local Similarity 58.6%; Pred. No. 0.00075;
Matches 99; Conservative 6; Mismatches 64; Indels 0; Gaps 0;

QY 3439 TTAGCGCAGTGTGCCAGGTGATGGATGATGACGACTACCGCGCAGAGATGACGATGAG 3498
DB 434 TTGSCDCAATGGTGGAGGCCATAGTGGAGTTTGATTACCAGGCCAGCATGATGATGAG 493
QY 3499 CTGCGCTTCAACAGAGGCCAGATCATCACTCTCTCAACAGAGGAGGCCCTGACTGTGG 3558
DB 494 CTGACGATCAGCGTGGGTGAGGTTCATCACCCWACCTCAGGATGAGGCTGGTGG 553
QY 3559 AAAGGAGAGTCAATGACAGAGTGGGCTCTTCCCATCCCAATTATGTGA 3607
DB 554 GAGGACAGATCAACGGCAGGAGAGGTTTGTTCCTGCACTTTGTAA 602
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RESULT 13
US-11-011-332A-156/c
; Sequence 156, Application US/11011332A
; Publication No. US20060024684A1
; GENERAL INFORMATION:
; APPLICANT: Roekens, John
; APPLICANT: Harbeck, Nadia
; APPLICANT: Koenig, Thomas
; APPLICANT: Maier, Sabine
; APPLICANT: Martens, John
; APPLICANT: Model, Fabian
; APPLICANT: Nimmrich, Inko
; APPLICANT: Rujan, Tamas
; APPLICANT: Schmitt, Manfred
; APPLICANT: Lesche, Ralf
; APPLICANT: Dietrich, Dima
; APPLICANT: Mueller, Volkmar
; APPLICANT: Kluth, Antje
; APPLICANT: Schwabe, Ina
; APPLICANT: Hartmann, Oliver
; APPLICANT: Adorjan, Peter
; TITLE OF INVENTION: PROGNOSTIC MARKERS FOR PREDICTION OF TREATMENT RESPONSE AND/OR SURVIVAL IN BREAST CANCER PATIENTS
; FILE REFERENCE: 47675-99
; CURRENT APPLICATION NUMBER: US/11/011,332A
; CURRENT FILING DATE: 2004-12-13
; PRIOR APPLICATION NUMBER: US 10/517,741
; PRIOR FILING DATE: 2003-10-01
; PRIOR APPLICATION NUMBER: PCT/EP2003/010881
; PRIOR FILING DATE: 2003-10-01
; PRIOR APPLICATION NUMBER: DE 10245779.4
; PRIOR FILING DATE: 2002-10-01
; PRIOR APPLICATION NUMBER: DE 10300096.8
; PRIOR FILING DATE: 2003-01-07
; PRIOR APPLICATION NUMBER: DE 10317955.0
; PRIOR FILING DATE: 2003-04-17
; PRIOR APPLICATION NUMBER: PCT/EP2004/014170
; PRIOR FILING DATE: 2004-12-13
; PRIOR APPLICATION NUMBER: EP 03090432.0
; PRIOR FILING DATE: 2003-12-11
; PRIOR APPLICATION NUMBER: EP 04090041.7
; PRIOR FILING DATE: 2004-02-10
; PRIOR APPLICATION NUMBER: EP 04090380.9
; PRIOR FILING DATE: 2004-09-30
; PRIOR APPLICATION NUMBER: EP 04090127.4
; PRIOR FILING DATE: 2004-04-01
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 158
; SEQ ID NO 156
; LENGTH: 28536
; TYPE: DNA
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: chemically treated genomic DNA (Homo sapiens)
US-11-011-332A-156

Query Match          1.2%; Score 63.8; DB 11; Length 28536;
Best Local Similarity 45.7%; Pred. No. 0.002;
Matches 273; Conservative 0; Mismatches 312; Indels 12; Gaps 1;

QY 1561 AATCGAGGCAAGCACTACTTAATCAAGAAACAAAGAAAGAGGACATAGTTGTACTG 1620
DB 6698 AATCAAAACCAAAACCAAAACCAAAACCAAAACCAAAACCAAAACCAAAACCAAAAC 6639
QY 1621 AAAGCAAGAAAAAGACTTTTGGATTTGAATTTGAAGCTCTTAATGATAAAAGCATCAA 1680
DB 6638 CAACCAAAACCAAAACCAAAACCAAAACCAAAACCAAAACCAAAACCAAAACCAAA 6579
QY 1681 CTGAAGAGGAACCTTCAAGATATCAGATGTGATGTGATGATGATGATGATGATGATG 1740
DB 6578 AATAAAAAAATAAAAAAATAAAAAAATAAAAAAATAAAAAAATAAAAAAATAAAAAA 6519
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1741 ACCACAAACAAATCTAGAGAGTTGAGAAATTGCCGAAATCACCCATCTACAGCAACAATTA 1800
6518 AACACCAAAACAAACAAACAAACAAACAAACAAACAAACAAACAAACAAACAAACAAACAA 6459
1801 CAGGAATCTCAGCAATGCTTGAAGACTTATTCAGAGAAACAGATCTCAATGACCA 1860
6458 AAAAAACAAACAAACAAACAAACAAACAAACAAACAAACAAACAAACAAACAAACAAATTA 6399
1861 TTAACAAAGTTTCAGCAGAACAGTTTGCACAGAGATTCTACTGTTACACTTAAAGAGCC 1920
6398 TTTAAACAAACAAACATATAAATATAAATATAAATATAAATATAAATATAAATATAAATATA 6339
1921 TTGAAGCAAAAGAACTAGCTCGGCAGCACTACGAGACCACTGGATGAAGTGGAGAA 1980
6338 TACAAAATTAACAAACAAACAAACAAACAAACAAACAAACAAACAAACAAACAAACAAACAA 6291
1981 GAACTAGATCAAACTACAGAGATTGATATTTCAATTAATCAGCTGAGGAACCTAAGA 2040
6290 AAAAAATAAAACCAATAAAACAAACAAACAAACAAACAAACAAACAAACAAACAAACAAAC 6231
2041 GAAATACACAATAAGCAACAACTCCAGAGCAAAAGTCCATGGAGGCTGAACGACTGAA 2100
6230 TAAATATAAAACAACTAAACAAACAAACAAACAAACAAACAAACAAACAAACAAACAAAC 6171
2101 CAGAAAGAAACAGAAAGATCTAGAAATTAGAAAAACAAAGAAAGAAAGCCCA 2157
6170 ACCAAAAACAAACAACTAACTTATCAACTAAAAACAAACAAACAAACAACTAAACCCCA 6114

RESULT 14

US-11-000-688-163
; Sequence 163, Application US/11000688
; Publication No. US20050287544A1
; GENERAL INFORMATION:
; APPLICANT: BERTUCCI, Francois
; APPLICANT: HOULGATTE, Remi
; TITLE OF INVENTION: GENE EXPRESSION PROFILING OF COLON CANCER WITH DNA ARRAYS
; FILE REFERENCE: 1423-R-03
; CURRENT APPLICATION NUMBER: US/11/000,688
; PRIOR FILING DATE: 2004-12-01
; PRIOR APPLICATION NUMBER: US 60/525,987
; NUMBER OF SEQ ID NOS: 1596
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 163
; LENGTH: 4257
; TYPE: DNA
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial sequences:primer
; NAME/KEY: misc feature
; LOCATION: (1)..(4257)
; OTHER INFORMATION: epidermal growth factor receptor pathway
; OTHER INFORMATION: substrate 15(EPS15) gene.
US-11-000-688-163

Query Match 1.2%; Score 63.2; DB 11; Length 4257;
Best Local Similarity 56.9%; Pred. No. 0.0011;
Matches 116; Conservative 0; Mismatches 88; Indels 0; Gaps 0;
250 TGGGCCATAACTGTAGAGAAAGCGAAGCATGATCAGCAGTTCATAGTTTAAAGCCA 309
456 TGGGCTGTAAACCTGAAGTAAGGCCAAATATGATGCAATATTGATAGTTTAAAGCCA 515
310 ATATCTGGATTCTACTGATCAGCTAGAAACTTTTTTTTCAATCTGGGTTACCT 369
516 GTGAATGGAAATTTCTGTCGTGATAAAGTGAACCAAGTGTGCTCAACTTAAGTTACCT 575
370 CAACCTGTTTATAGCAGAGATATGGCACTAGCTGACATGAATAATGATGAAGAATGGAT 429

576 GTGGATATCCTTTGGAGAGATTGGAGTTGAGTATGATATGACCATGATGGAATGCTTGAC 635
430 CAAGTGGAGTTTTCATAGCTATG 453
636 AGAGATGAGTTTGCAAGTTGCCATG 659
RESULT 15
US-11-000-688-1278
; Sequence 1278, Application US/11000688
; Publication No. US20050287544A1
; GENERAL INFORMATION:
; APPLICANT: BERTUCCI, Francois
; APPLICANT: HOULGATTE, Remi
; APPLICANT: BIRNBAUM, Daniel
; TITLE OF INVENTION: GENE EXPRESSION PROFILING OF COLON CANCER WITH DNA ARRAYS
; FILE REFERENCE: 1423-R-03
; CURRENT APPLICATION NUMBER: US/11/000,688
; PRIOR FILING DATE: 2004-12-01
; PRIOR APPLICATION NUMBER: US 60/525,987
; NUMBER OF SEQ ID NOS: 1596
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 1278
; LENGTH: 2774
; TYPE: DNA
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial sequences:primer
; NAME/KEY: misc feature
; LOCATION: (1)..(2774)
; OTHER INFORMATION: epidermal growth factor receptor substrate
; OTHER INFORMATION: eps15r(EPS15R) gene.
US-11-000-688-1278

Query Match 1.2%; Score 62.8; DB 11; Length 2774;
Best Local Similarity 52.3%; Pred. No. 0.0011;
Matches 139; Conservative 0; Mismatches 127; Indels 0; Gaps 0;
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366 CTGGGCTGTAGGTTGGAGAAAGGCCAAATTTGATGGGATTTTGAAGCCTCTTGCC 425
309 AATATCTGGATTCTACTGATCAAGCTAGAAACTTTTTTTTCAATCTGGGTTACC 368
426 CATCAATGGTTGCTCTCTGGAGACAAAGTCAAGCCAGTCTCATGAACCTCAAGCTGCC 485
369 TCAACCTGTTTATAGCAGAGATATGGCACTAGCTGACATGAATATGATGAAGAATGGA 428
486 TCTTGATGCTCTGGGACGGTCTGGACCTCAGTGACATTGACAAGGATGGGCACTTGA 545
429 TCAAGTGGAGTTTTCATAGCTATGAAACTTATCAAACTGAAAGCTACAAGGATATCAGCT 488
546 TCGAGATGAGTTCGCTGTGGCCATGCACTTGGTGTACCGAGCCCTGGAGAGAGCCCT 605
489 ACCCTCTGCACCTTCCCTCTGTCATGA 514
606 GCCCTCCGCCCTGCCCGCTCCCTCA 631

Search completed: February 14, 2006, 05:59:35
Job time : 752 secs

GenCore version 5.1.7
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OM nucleic - nucleic search, using sw model

Run on: February 14, 2006, 05:22:49 ; Search time 3772 Seconds
(without alignments)
11397.799 Million cell updates/sec

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Perfect score: 5199
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Scoring table: IDENTITY NUC
Gapop 10.0 , Gapext 1.0

Searched: 9793542 seqs, 4134689005 residues

Total number of hits satisfying chosen parameters: 19587084

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Published Applications NA.Main:*
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3: /cgn2_6/ptodata/1/pubpna/US09A_PUBCOMB.seq:*
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9: /cgn2_6/ptodata/1/pubpna/US10E_PUBCOMB.seq:*
10: /cgn2_6/ptodata/1/pubpna/US11_PUBCOMB.seq:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

| Result No. | Score | Query Match % | Length | DB ID | Description |
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| 1 | 3285.6 | 63.2 | 7247 | 8 | US-10-852-943-84 |
| 2 | 3285.6 | 63.2 | 7247 | 9 | US-10-287-436A-57 |
| 3 | 3285.6 | 63.2 | 7247 | 9 | US-10-287-436A-685 |
| 4 | 3090.6 | 59.4 | 7435 | 9 | US-10-450-763-20567 |
| 5 | 2884.8 | 55.5 | 3466 | 7 | US-10-158-057-33 |
| 6 | 2843.2 | 54.7 | 3319 | 6 | US-09-764-875-88 |
| 7 | 1882.4 | 36.2 | 2067 | 6 | US-10-264-049-887 |
| 8 | 1630 | 31.4 | 2874 | 9 | US-10-450-763-20566 |
| 9 | 550 | 10.6 | 5828 | 7 | US-10-398-885A-15 |
| 10 | 507.8 | 9.8 | 2017 | 3 | US-09-884-441-72 |
| 11 | 507.8 | 9.8 | 2017 | 3 | US-09-907-969-72 |
| 12 | 507.8 | 9.8 | 2017 | 3 | US-09-827-271-72 |
| 13 | 507.8 | 9.8 | 2017 | 6 | US-10-198-053-72 |
| 14 | 507.8 | 9.8 | 2017 | 6 | US-10-860-790-72 |
| 15 | 503.6 | 9.7 | 568 | 3 | US-09-764-881-55 |
| 16 | 503.6 | 9.7 | 568 | 3 | US-09-764-881-55 |
| 17 | 503.6 | 9.7 | 568 | 3 | US-09-764-875-404 |
| 18 | 503.6 | 9.7 | 568 | 6 | US-10-242-747-55 |
| 19 | 503.6 | 9.7 | 568 | 7 | US-10-158-057-127 |
| 20 | 462.6 | 8.9 | 503 | 3 | US-09-918-995-31258 |
| 21 | 291.4 | 5.6 | 395 | 9 | US-10-450-763-20563 |
| 22 | 264.2 | 5.1 | 301 | 3 | US-09-864-761-17146 |
| 23 | 263 | 5.1 | 263 | 3 | US-09-864-761-17644, A |

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| 24 | 250 | 4.8 | 270 | 3 | US-09-864-761-17127 | Sequence 17127, A |
| 25 | 250 | 4.8 | 286 | 3 | US-09-864-761-26948 | Sequence 26948, A |
| 26 | 250 | 4.8 | 297 | 3 | US-09-864-761-30453 | Sequence 30453, A |
| 27 | 240 | 4.6 | 486 | 3 | US-09-864-761-333 | Sequence 333, App |
| 28 | 240 | 4.6 | 487 | 3 | US-09-864-761-864 | Sequence 864, App |
| 29 | 239 | 4.6 | 480 | 3 | US-09-864-761-10314 | Sequence 10314, A |
| 30 | 237.8 | 4.6 | 418 | 3 | US-09-783-590-8760 | Sequence 8760, App |
| 31 | 230 | 4.4 | 230 | 3 | US-09-864-761-17643 | Sequence 17643, A |
| 32 | 230 | 4.4 | 247 | 3 | US-09-864-761-30501 | Sequence 30501, A |
| 33 | 205 | 3.9 | 484 | 3 | US-09-864-761-863 | Sequence 863, App |
| 34 | 202 | 3.9 | 475 | 3 | US-09-864-761-311 | Sequence 311, App |
| 35 | 202 | 3.9 | 475 | 3 | US-09-864-761-13884 | Sequence 13884, A |
| 36 | 186 | 3.6 | 477 | 3 | US-09-864-761-13936 | Sequence 13936, A |
| 37 | 180 | 3.5 | 180 | 3 | US-09-864-761-17125 | Sequence 17125, A |
| 38 | 167.6 | 3.2 | 3981 | 10 | US-11-097-143-233 | Sequence 233, App |
| 39 | 165.6 | 3.2 | 967 | 3 | US-09-764-881-50 | Sequence 50, Appl |
| 40 | 165.6 | 3.2 | 967 | 3 | US-09-764-881-50 | Sequence 50, Appl |
| 41 | 165.6 | 3.2 | 967 | 3 | US-09-764-875-411 | Sequence 411, App |
| 42 | 165.6 | 3.2 | 967 | 6 | US-10-242-747-50 | Sequence 50, Appl |
| 43 | 165.6 | 3.2 | 967 | 7 | US-10-158-057-40 | Sequence 40, Appl |
| 44 | 165.4 | 3.2 | 292 | 3 | US-09-864-761-20261 | Sequence 20261, A |
| 45 | 165.4 | 3.2 | 304 | 3 | US-09-864-761-21373 | Sequence 21373, A |

ALIGNMENTS

RESULT 1
US-10-852-943-84
; Sequence 84, Application US/10852943
; Publication No. US20050037388A1
; GENERAL INFORMATION:
; APPLICANT: University of Geneva
; APPLICANT: Stylianos, Antonarakis
; APPLICANT: Deutsch, Samuel
; TITLE OF INVENTION: METHOD FOR DETECTING DISEASES CAUSED BY CHROMOSOMAL IMBALANCES
; FILE REFERENCE: 27067/2005
; CURRENT APPLICATION NUMBER: US/10/852,943
; CURRENT FILING DATE: 2004-05-25
; PRIOR APPLICATION NUMBER: US 60/300,266
; PRIOR FILING DATE: 2001-06-21
; PRIOR APPLICATION NUMBER: US 10/177,063
; PRIOR FILING DATE: 2002-06-21
; NUMBER OF SEQ ID NOS: 98
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 84
; LENGTH: 7247
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-852-943-84

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| Query Match | 63.2% | Score | 3285.6 | DB 8 | Length | 7247 | | | |
| Best Local Similarity | 93.8% | Pred. No. | 0 | | | | | | |
| Matches | 3536 | Conservative | 0 | Mismatches | 4 | Indels | 228 | Gaps | 2 |

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| Qy | 102 | GCCTCCCTCCACGGCGCTGAGCGCACTGATTGTCCTCGGGCGGCGCGGACG | 161 |
| Db | 1 | CGCTCCCTCCACGGCGCTGAGCGCACTGATTGTCCTCGGGCGGCGCGGACG | 60 |
| Qy | 162 | CGCCCGGAGATGAGCGCTCGATTAGCAAGTAAAGTAAACAGAACCATGGCTCAGTTTC | 221 |
| Db | 61 | CGCCCGGAGATGAGCGCTCGATTAGCAAGTAAAGTAAACAGAACCATGGCTCAGTTTC | 120 |
| Qy | 222 | AACACCTTTGGTGGCAGCTCGATATCTGGCCATACCTGTAGAGAAAGAGCGGAGCA | 281 |
| Db | 121 | AACACCTTTGGTGGCAGCTCGATATCTGGCCATACCTGTAGAGAAAGAGCGGAGCA | 180 |
| Qy | 282 | TGATCAGCAGTTCATAGTTTAAAGCCAATATCTGGATTCTGATTCAGTCAAGCTAG | 341 |
| Db | 181 | TGATCAGCAGTTCATAGTTTAAAGCCAATATCTGGATTCTGATTCAGTCAAGCTAG | 240 |
| Qy | 342 | AAACCTTTTTTTCAATCTCGGTTACTCAACCTGTTTATAGCAGATATGGGCACATGC | 401 |

| | | | |
|----|------|---|------|
| Db | 241 | AAACTTTTTTTTCAATCTGGGTTACCTCAACTGTTTTAGCACAGATATGGGCACTAGC | 300 |
| Qy | 402 | TGACATGAATAATGATGGAAGAATGGATCAAGTGGAGTTTTCCATAGCTATGAAACTTAT | 461 |
| Db | 301 | TGACATGAATAATGATGGAAGAATGGATCAAGTGGAGTTTTCCATAGCTATGAAACTTAT | 360 |
| Qy | 462 | CAAACTGAAGCTACAAGGATATCAGCTACCTCTGCACTTCCCTCTGTCATGAAGACGA | 521 |
| Db | 361 | CAAACTGAAGCTACAAGGATATCAGCTACCTCTGCACTTCCCTCTGTCATGAAGACGA | 420 |
| Qy | 522 | ACCAGTTGCTATTTCTAGCGCACAGCATTTGGTATGGAGGTATCGCCAGCATGCCACC | 581 |
| Db | 421 | ACCAGTTGCTATTTCTAGCGCACAGCATTTGGTATGGAGGTATCGCCAGCATGCCACC | 480 |
| Qy | 582 | GCTTACAGCTGTGTCTCCAGTCCCAATGGGATCCATTCAGTGTGTGGAAATGCTCCAAC | 641 |
| Db | 481 | GCTTACAGCTGTGTCTCCAGTCCCAATGGGATCCATTCAGTGTGTGGAAATGCTCCAAC | 540 |
| Qy | 642 | CCTAGTATCTTCTGTTCACAGCAGCTGTGCCCCCTGGCTTAAGGGGCTCCCCCTGT | 701 |
| Db | 541 | CCTAGTATCTTCTGTTCACAGCAGCTGTGCCCCCTGGCTTAAGGGGCTCCCCCTGT | 600 |
| Qy | 702 | TATACAACTCTGCTGCAATTTGCTCATCTGCAGCACATTTGCCAAAGAGTTCTTCCCT | 761 |
| Db | 601 | TATACAACTCTGCTGCAATTTGCTCATCTGCAGCACATTTGCCAAAGAGTTCTTCCCT | 660 |
| Qy | 762 | TAGTAGATCTGCTCCAGGGTCAACAATAAACACTAAATTACAAAAGGCAAGTCAATTTGA | 821 |
| Db | 661 | TAGTAGATCTGCTCCAGGGTCAACAATAAACACTAAATTACAAAAGGCAAGTCAATTTGA | 720 |
| Qy | 822 | TGTGGCAGTGTCCACAGTGGCAGAGTGGGCTGTTCTCAGTCATCAAGACTGAATA | 881 |
| Db | 721 | TGTGGCAGTGTCCACAGTGGCAGAGTGGGCTGTTCTCAGTCATCAAGACTGAATA | 780 |
| Qy | 882 | CAGGCAATTTATTTCAATAGTCATGACAAAATATAGTGGACACTTAAACAGTCCCCAAGC | 941 |
| Db | 781 | CAGGCAATTTATTTCAATAGTCATGACAAAATATAGTGGACACTTAAACAGTCCCCAAGC | 840 |
| Qy | 942 | AAGAACTATTTTATGCACTCAAGTTTACCACAGGCTCAGCTGGCTTCAATATGGAATCT | 1001 |
| Db | 841 | AAGAACTATTTTATGCACTCAAGTTTACCACAGGCTCAGCTGGCTTCAATATGGAATCT | 900 |
| Qy | 1002 | TTCTGACATGTACAGATGGAAGAACTTACAGCAGAGGAATTTATCCTGGCAATGCACCT | 1061 |
| Db | 901 | TTCTGACATGTACAGATGGAAGAACTTACAGCAGAGGAATTTATCCTGGCAATGCACCT | 960 |
| Qy | 1062 | CATTGATGTAGTATGCTCTGGCCAAACCACTGCCACCTGCTCCTGCCCTCCAGAAATACCTCC | 1121 |
| Db | 961 | CATTGATGTAGTATGCTCTGGCCAAACCACTGCCACCTGCTCCTGCCCTCCAGAAATACCTCC | 1020 |
| Qy | 1122 | ACCTTCTTTTGAAGAGTTTCAATCTGGCAGTGGTATATCTGTCTATAAGCTCAACATCTGT | 1181 |
| Db | 1021 | ACCTTCTTTTGAAGAGTTTCAATCTGGCAGTGGTATATCTGTCTATAAGCTCAACATCTGT | 1080 |
| Qy | 1182 | AGATCAGAGCTTACAGAGGACCACTTTTAGNAGTGAACAACAACAAATTAGAAGAA | 1241 |
| Db | 1081 | AGATCAGAGCTTACAGAGGACCACTTTTAGAAGATGAACAACAACAATTAGAAGAA | 1140 |
| Qy | 1242 | ATTACTGTAAACGTTTGAAGATAAGAGCGGAGAACTTTGAACCTGGCAACCTGGAACCT | 1301 |
| Db | 1141 | ATTACTGTAAACGTTTGAAGATAAGAGCGGAGAACTTTGAACCTGGCAACCTGGAACCT | 1200 |
| Qy | 1302 | GGAGAAACGAAGGCAAGCTCTCTCGAAACAGCAGCGCAAGGAGCAGGAGCGCTGGCCCA | 1361 |
| Db | 1201 | GGAGAAACGAAGGCAAGCTCTCTCGAAACAGCAGCGCAAGGAGCAGGAGCGCTGGCCCA | 1260 |
| Qy | 1362 | GCTGGAGCGGGCGGAGCAGAGAGGAGAGCGGTGAGCCCGCAGGAGCAGAGCGCAAGAG | 1421 |
| Db | 1261 | GCTGGAGCGGGCGGAGCAGAGAGGAGAGCGGTGAGCCCGCAGGAGCAGAGCGCAAGAG | 1320 |
| Qy | 1422 | ACAACTGGAACTGGAGAACCAACTGGAAAAGCAGCGGGAGCTTAGAACCGCAGAGAGGA | 1481 |
| Db | 1321 | ACAACTGGAACTGGAGAACCAACTGGAAAAGCAGCGGGAGCTTAGAACCGCAGAGAGGA | 1380 |
| Qy | 1482 | GGAGAGGAGGAAAGAAATTTGAGAGCGGAGAGGCTGCAAAAACGGGAACTTTGAAAGCAACG | 1541 |
| Db | 1381 | GGAGAGGAGGAAAGAAATTTGAGAGCGGAGAGGCTGCAAAAACGGGAACTTTGAAAGCAACG | 1440 |
| Qy | 1542 | ACAACTTCAGTGGGAACGGAAATCGAAGCAAGAACTACTAAATCAAGAAACAAAGAAACA | 1601 |
| Db | 1441 | ACAACTTCAGTGGGAACGGAAATCGAAGCAAGAACTACTAAATCAAGAAACAAAGAAACA | 1500 |
| Qy | 1602 | AGAGGACATAGTTGTACTGAAAGCAAAAGAACTTTTGGAAATTTGAAATAGAAAGCTCT | 1661 |
| Db | 1501 | AGAGGACATAGTTGTACTGAAAGCAAAAGAACTTTTGGAAATTTGAAATAGAAAGCTCT | 1560 |
| Qy | 1662 | AAATGATATAAAGCATTCACTAGAAAGGAACTTCAAGATATCAGATGTCGATTCACCCAC | 1721 |
| Db | 1561 | AAATGATATAAAGCATTCACTAGAAAGGAACTTCAAGATATCAGATGTCGATTCACCCAC | 1620 |
| Qy | 1722 | CCAAAGGCAAGAAATTTGAGAGCACAAACAAATCTAGAGAGTTTGAAATTTGCCGAATCAC | 1781 |
| Db | 1621 | CCAAAGGCAAGAAATTTGAGAGCACAAACAAATCTAGAGAGTTTGAAATTTGCCGAATCAC | 1680 |
| Qy | 1782 | CCATCTACAGCAACAATTTACAGGAATCTCAGCAAAATGCTTGGAGACTTATTTCCAGAAA | 1841 |
| Db | 1681 | CCATCTACAGCAACAATTTACAGGAATCTCAGCAAAATGCTTGGAGACTTATTTCCAGAAA | 1740 |
| Qy | 1842 | ACAGATCTCAATGACCAATTTAAACAAGTTTCAGCAGAACAGTTTGGCAGAGATTCAT | 1901 |
| Db | 1741 | ACAGATCTCAATGACCAATTTAAACAAGTTTCAGCAGAACAGTTTGGCAGAGATTCAT | 1800 |
| Qy | 1902 | TGTTACACTTAAAGAGGCTTTAGAAAGCAAAAGAACTAGCTCGGAGACCTTACAGAGACA | 1961 |
| Db | 1801 | TGTTACACTTAAAGAGGCTTTAGAAAGCAAAAGAACTAGCTCGGAGACCTTACAGAGACA | 1860 |
| Qy | 1962 | ACTGATGAAGTGGAGAAAGAACTTAGATCAAAACTACAGAGATTTGATATTTTCAATAA | 2021 |
| Db | 1861 | ACTGATGAAGTGGAGAAAGAACTTAGATCAAAACTACAGAGATTTGATATTTTCAATAA | 1920 |
| Qy | 2022 | TCAGCTGAAGAACTTAAGAGAAATACAAATAAGCAACAATCCAGAGAGCAAAAGTCCAT | 2081 |
| Db | 1921 | TCAGCTGAAGAACTTAAGAGAAATACAAATAAGCAACAATCCAGAGAGCAAAAGTCCAT | 1980 |
| Qy | 2082 | GGAGGCTCAACGACTGAAACAGAAAGAAACAAAGAACTATAGAAATTTAGAAAAACA | 2141 |
| Db | 1981 | GGAGGCTCAACGACTGAAACAGAAAGAAACAAAGAACTATAGAAATTTAGAAAAACA | 2040 |
| Qy | 2142 | AAAAGAAAGCCCAAGACGAGCTCAGGAAAGGCAAGCAGTGGCTGGAGCATGTGCA | 2201 |
| Db | 2041 | AAAAGAAAGCCCAAGACGAGCTCAGGAAAGGCAAGCAGTGGCTGGAGCATGTGCA | 2100 |
| Qy | 2202 | GCAGGAGCAGCATCAGAGACCAAGAAACTCCACCAAGGAAACCTGAAAAAGGGA | 2261 |
| Db | 2101 | GCAGGAGCAGCATCAGAGACCAAGAAACTCCACCAAGGAAACCTGAAAAAGGGA | 2160 |
| Qy | 2262 | GGAGAGTCTCAAAAAGAGGATGGCGAGAAAAAGGCAAAACAGGAAGCACAAGACAAGCT | 2321 |
| Db | 2161 | GGAGAGTCTCAAAAAGAGGATGGCGAGAAAAAGGCAAAACAGGAAGCACAAGACAAGCT | 2220 |
| Qy | 2322 | GGGTCGGCTTTTCCATCAACCAAGAACAGCTTAAGCCAGCTGTCCAGGCAACCTCGTTC | 2381 |
| Db | 2221 | GGGTCGGCTTTTCCATCAACCAAGAACAGCTTAAGCCAGCTGTCCAGGCAACCTCGTTC | 2280 |
| Qy | 2382 | CAGTCAGAAAAAGGTCACCTTACCTTCTGCAAGGAAATGTAAGTGGTCTATTATA | 2441 |
| Db | 2281 | CAGTCAGAAAAAGGTCACCTTACCTTCTGCAAGGAAATGTAAGTGGTCTATTATA | 2340 |
| Qy | 2442 | CGGGCACTGTACCCCTTTTGAATCCAGAAAGCCATGATGAAATCACTATCCAGCCAGGAGA | 2501 |
| Db | 2341 | CGGGCACTGTACCCCTTTTGAATCCAGAAAGCCATGATGAAATCACTATCCAGCCAGGAGA | 2400 |
| Qy | 2502 | CATAGTCTAT-----GGTGGATGAAAGCCAAACTGAGAGAACCCCGCTGGCT | 2546 |
| Db | 2401 | CATAGTCTATGTTAAAGGGGAATGGGTGGATGAAAGCCAAACTGAGAGAACCCCGCTGGCT | 2460 |

QY 582 GCTTACAGCTGTGCTCCAGTGCCAAATGGGATCCATTCCAGTTGTTGGAATGTCCTCAAC 641
DB 481 GCTTACAGCTGTGCTCCAGTGCCAAATGGGATCCATTCCAGTTGTTGGAATGTCCTCAAC 540
QY 642 CCTAGTATCTCTGTTCCACAGCAGCTGTGCCCCCTGGCTTAACGGGGCTCCCCCTGT 701
DB 541 CCTAGTATCTCTGTTCCACAGCAGCTGTGCCCCCTGGCTTAACGGGGCTCCCCCTGT 600
QY 702 TATACAACTCTGCTGCAATTTGCTCATCTCGAGCCACATTTGCCAAAGAGTTCTTCCTT 761
DB 601 TATACAACTCTGCTGCAATTTGCTCATCTCGAGCCACATTTGCCAAAGAGTTCTTCCTT 660
QY 762 TAGTAGATCTGCTCCAGGGTCACTAAACACATAAATTTACAAAGGCACAGTCATTGA 821
DB 661 TAGTAGATCTGCTCCAGGGTCACTAAACACATAAATTTACAAAGGCACAGTCATTGA 720
QY 822 TGTGGCCAGTGTCCACACAGTGGCAGAGTGGGCTGTTCTCAGTCATCAAGACTGAATA 881
DB 721 TGTGGCCAGTGTCCACACAGTGGCAGAGTGGGCTGTTCTCAGTCATCAAGGCTGAATA 780
QY 882 CAGGCCAATTAATCAATAGTCATGACAAAACTATGAGTGGACACTTAAACAGGTCCTCAAGC 941
DB 781 CAGGCCAATTAATCAATAGTCATGACAAAACTATGAGTGGACACTTAAACAGGTCCTCAAGC 840
QY 942 AAGAACTATTTCTATGCACTCAAGTTTACACAGGCTCAGCTGGCTTCAATATGGAATCT 1001
DB 841 AAGAACTATTTCTATGCACTCAAGTTTACACAGGCTCAGCTGGCTTCAATATGGAATCT 900
QY 1002 TTCTGACATTTGATCAAGATGGAATACTTACAGCAGAGGAATTTATCCTGGCAATGCACCT 1061
DB 901 TTCTGACATTTGATCAAGATGGAATACTTACAGCAGAGGAATTTATCCTGGCAATGCACCT 960
QY 1062 CATTTGATGTAGTATGTCTGGCCAACTCACTGGCACCTGTCTGCTCCAGAAATACATTC 1121
DB 961 CATTTGATGTAGTATGTCTGGCCAACTCACTGGCACCTGTCTGCTCCAGAAATACATTC 1020
QY 1122 ACCTTCTTTTGAAGAGTTGATCTGGCAGTGGTATATCTGTCTCAATAGCTCAACATCTGT 1181
DB 1021 ACCTTCTTTTGAAGAGTTGATCTGGCAGTGGTATATCTGTCTCAATAGCTCAACATCTGT 1080
QY 1182 AGATCAGAGGCTACCAAGAGAACCCAGTTTTAGAAGATGAACAACTTAGAAAGAA 1241
DB 1081 AGATCAGAGGCTACCAAGAGAACCCAGTTTTAGAAGATGAACAACTTAGAAAGAA 1140
QY 1242 ATTACCTGTAACTTTGAAGATAAGAACGGGAGAACTTTGAACGTGGCAACTGGAAC 1301
DB 1141 ATTACCTGTAACTTTGAAGATAAGAACGGGAGAACTTTGAACGTGGCAACTGGAAC 1200
QY 1302 GGAGAAACGAAGGCAAGCTCTCTGGAAACAGCAGCGCAAGGACAGGAGCGCTGGCCCA 1361
DB 1201 GGAGAAACGAAGGCAAGCTCTCTGGAAACAGCAGCGCAAGGACAGGAGCGCTGGCCCA 1260
QY 1362 GCTGGAGCGGGCGGAGCAGGAGGAAGGAGCGTGAGCGCCAGGAGCAGCGCAAG 1421
DB 1261 GCTGGAGCGGGCGGAGCAGGAGGAAGGAGCGTGAGCGCCAGGAGCAGCGCAAG 1320
QY 1422 ACAACTGGAACTGGGAAAGCAACTGGAAGCAGCGGGAGCTAGAACCGCAGAGAGGA 1481
DB 1321 ACAACTGGAACTGGGAAAGCAACTGGAAGCAGCGGGAGCTAGAACCGCAGAGAGGA 1380
QY 1482 GGAGAGGAGGAAGAAATTTGAGAGCGGAGAGCGTGCAAAACGGGAACTTTGAAAGGCAAG 1541
DB 1381 GGAGAGGAGGAAGAAATTTGAGAGCGGAGAGCGTGCAAAACGGGAACTTTGAAAGGCAAG 1440
QY 1542 ACAACTTTGAGTGGGAACGAATTCGAAGGCAAGAACTTACTTAATCAAGAAAAACAAGAA 1601
DB 1441 ACAACTTTGAGTGGGAACGAATTCGAAGGCAAGAACTTACTTAATCAAGAAAAACAAGAA 1500
QY 1602 AGAGGACATAGTTGTACTGAAAGCAAGAAAAAGACTTTTGGAAATTTGAAATTTGAAGCTCT 1661
DB 1501 AGAGGACATAGTTGTACTGAAAGCAAGAAAAAGACTTTTGGAAATTTGAAATTTGAAGCTCT 1560
QY 1662 AATGATAAAGACATCACTAGAAAGGGAACTTCAAGATATCAGATGTCGATTGACCC 1721

DB 1561 AAATGATAAAGACATCACTAGAAAGGAAACTTCAAGATATCAGATGTCGATTGACCAC 1620
QY 1722 CCAAAGGCAGAAATTTGAGAGCACAAAACAAATCTAGAGAGTTGAGAAATTCGCCAAATCAC 1781
DB 1621 CCAAAGGCAGAAATTTGAGAGCACAAAACAAATCTAGAGAGTTGAGAAATTCGCCAAATCAC 1680
QY 1782 CCATCTCAGCAACAAATTTACAGGAATCTCAGCAAAATGCTTGGAGACATTTATTCAGAAAA 1841
DB 1681 CCATCTCAGCAACAAATTTACAGGAATCTCAGCAAAATGCTTGGAAAGACTTATTTCCAGAAAA 1740
QY 1842 ACAGATATCTCAATGACCAATTTAAACAGTTTCAGCAGAACTAGTTTGCACAGAGATTCACT 1901
DB 1741 ACAGATATCTCAATGACCAATTTAAACAGTTTCAGCAGAACTAGTTTGCACAGAGATTCACT 1800
QY 1902 TGTTACACTTAAAGAGCGCTTTAGAAAGCAAAAGAACTAGCTCGGCAGCACCTACCGAGCCA 1961
DB 1801 TGTTACACTTAAAGAGCGCTTTAGAAAGCAAAAGAACTAGCTCGGCAGCACCTACCGAGCCA 1860
QY 1962 ACTGATGAAAGTGGAGAAAGAACTAGATCAAACTACAGGAGATTGATTTTCAATAA 2021
DB 1861 ACTGATGAAAGTGGAGAAAGAACTAGATCAAACTACAGGAGATTGATTTTCAATAA 1920
QY 2022 TCAGCTGAAGAACTTAAGAGAAATACACAATAAGCAAACTCCAGAAAGCAAAAGTCCAT 2081
DB 1921 TCAGCTGAAGAACTTAAGAGAAATACACAATAAGCAAACTCCAGAAAGCAAAAGTCCAT 1980
QY 2082 GGAGGCTGAAACGACTGAAACAGAAAGAAACAAAGAACGAAAGATCATAGAAATTTAGAAAAACA 2141
DB 1981 GGAGGCTGAAACGACTGAAACAGAAAGAAACAAAGAACGAAAGATCATAGAAATTTAGAAAAACA 2040
QY 2142 AAAAGAAAGCCCAAAGACGAGCTCAGGAAAGGGAACGAGTGGCTGGAGCATGTGCA 2201
DB 2041 AAAAGAAAGCCCAAAGACGAGCTCAGGAAAGGGAACGAGTGGCTGGAGCATGTGCA 2100
QY 2202 CGAGGAGCAGGATCAGAGCAAGAAACTCCAGAAAGGAAACTTGAAGAAAGGA 2261
DB 2101 CGAGGAGCAGGATCAGAGCAAGAAACTCCAGAAAGGAAACTTGAAGAAAGGA 2160
QY 2262 GGAGAGTGTCAAAAGAGGATGGCGAGAAAAAGGCAAAACGAGGACCAAGACAAGCT 2321
DB 2161 GGAGAGTGTCAAAAGAGGATGGCGAGAAAAAGGCAAAACGAGGACCAAGACAAGCT 2220
QY 2322 GGGTCGGCTTTTCAATCAACCAAGAACAGCTTAAGCAGCTGTCCAGGACACCTGGTC 2381
DB 2221 GGGTCGGCTTTTCAATCAACCAAGAACAGCTTAAGCAGCTGTCCAGGACACCTGGTC 2280
QY 2382 CACTGCAAAAAAGGTCCTTACCATTCTGCAACAGGAAATGTAAAAGTGGTGATTA 2441
DB 2281 CACTGCAAAAAAGGTCCTTACCATTCTGCAACAGGAAATGTAAAAGTGGTGATTA 2340
QY 2442 CCGGCACTGTACCCCTTTGAAATCCAGAGCCATGATGAAATCACTATCCAGCCAGGAGA 2501
DB 2341 CCGGCACTGTACCCCTTTGAAATCCAGAGCCATGATGAAATCACTATCCAGCCAGGAGA 2400
QY 2502 CATAGTCAT-----GGTGGATGAAAGCCAAAACCTGGAGAACCCGGCTGGCT 2546
DB 2401 CATAGTCATGTTTAAAGGGGAATGGGTGGATGAAAGCCAAAACCTGGAGAACCCGGCTGGCT 2460
QY 2547 TGGAGGAAATTTAAAGGAAAGACAGGCTGGTCCCTGCAAACTATGCAGAGAAATCCC 2606
DB 2461 TGGAGGAAATTTAAAGGAAAGACAGGCTGGTCCCTGCAAACTATGCAGAGAAATCCC 2520
QY 2607 AGAAATGAGGTTCCCGCTCCAGTGAAACCAAGTGAATTTCAACATCTGCCCTGCCCC 2666
DB 2521 AGAAATGAGGTTCCCGCTCCAGTGAAACCAAGTGAATTTCAACATCTGCCCTGCCCC 2580
QY 2667 CAATGCGCTTGGTGAGACCCCGCCCTTTGGCAGTAACCTTTTCAGAGCCCTCCAC 2726
DB 2581 CAATGCGCTTGGTGAGACCCCGCCCTTTGGCAGTAACCTTTTCAGAGCCCTCCAC 2640
QY 2727 GACCCCTAATAACTTGGGCGGACTTCAGCTCCACGTGGCCCAACAGCAGCAATGAGAAACC 2786

Db 2641 GACCCCTAATAACTGGGCGCACTTCAGCTCCAGTGGGCCACAGCAGCAATGAGAAACC 2700
QY 2787 AGAAACGGATAACTGGGATGCAATGGGCGAGCCAGCCCTCTCTCACCGTTCCAAAGTCCCGG 2846
Db 2701 AGAAACGGATAACTGGGATGCAATGGGCGAGCCAGCCCTCTCTCACCGTTCCAAAGTCCCGG 2760
QY 2847 CCAGTTAAGGCGAGGTCGCGCTTTACTCCAGCCAGCGGCACTGGCTCTCCCGTCTCC 2906
Db 2761 CCAGTTAAGGCGAGGTCGCGCTTTACTCCAGCCAGCGGCACTGGCTCTCCCGTCTCC 2820
QY 2907 TGTGCTAGGCCAGGTTGAAAAGGTGAGGGGCTTACAAGCTCAAGCCCTATATCTTTGGAG 2966
Db 2821 TGTGCTAGGCCAGGTTGAAAAGGTGAGGGGCTTACAAGCTCAAGCCCTATATCTTTGGAG 2880
QY 2967 AGCCAAAAAGCAACCACTTAATTTTAAACAAAATGATGTCATCACGTCCTGGAACA 3026
Db 2881 AGCCAAAAAGCAACCACTTAATTTTAAACAAAATGATGTCATCACGTCCTGGAACA 2940
QY 3027 GCAAGACATGTGTGTTGGAGAGTTCAGGTCAAGGTGAGAGGGTGTGGTCCCAAGTCTTA 3086
Db 2941 GCAAGACATGTGTGTTGGAGAGTTCAGGTCAAGGTGAGAGGGTGTGGTCCCAAGTCTTA 3000
QY 3087 CGTGAACCTCAATTCAGGGGCCATAAGGAAGTCTACAAGCATGGAATTCGTTCCTCAGA 3146
Db 3001 CGTGAACCTCAATTCAGGGGCCATAAGGAAGTCTACAAGCATGGAATTCGTTCCTCAGA 3060
QY 3147 GAGTCTGTAGTCTAAAGGAGTAGCTCTCAGCAGCAAGCCGGTGTTCGGGAGA 3206
Db 3061 GAGTCTGTAGTCTAAAGGAGTAGCTCTCAGCAGCAAGCCGGTGTTCGGGAGA 3120
QY 3207 A----- 3207
Db 3121 AGAATTTATGTCATGTACACTTACGAGAGTTCTGAGCAAGGAGATTTAACCTTTTCAGCA 3180
QY 3208 ----- 3207
Db 3181 AGGGGATGTATTTGGTTTACCAGAAAGATGTGTGACTGTGTGGACAGGAACAGTGGGCGA 3240
QY 3208 ----- 3207
Db 3241 CAAGGCGGAGTCTTCCCTTCTAATATGTGAGGCTTAAAGATTACAGAGGCTCTGGAAC 3300
QY 3208 -----GAAATTCGCCAGGTATTGCTTCATA 3233
Db 3301 TGTGCGAAACAGGAGTTTAGAAAAAACCTGAAATTTGCCAGGTTATGCTTCATA 3360
QY 3234 CACCGCCACCGGCCCCGAGCAGCTCACTCTCGCCCTCTGTGTCAGCTGATTTGATCCGAAA 3293
Db 3361 CACCGCCACCGGCCCCGAGCAGCTCACTCTCGCCCTCTGTGTCAGCTGATTTGATCCGAAA 3420
QY 3294 AAAGAACCCAGGTGGATGTTGGGAAAGAGAGCTGCAAGCAAGTGGGAAAGAGCCGAGAT 3353
Db 3421 AAAGAACCCAGGTGGATGTTGGGAAAGAGAGCTGCAAGCAAGTGGGAAAGAGCCGAGAT 3480
QY 3354 AGGCTGGTCCAGCAATTTATGTAAAGCTTCTAAGCCCTGGGACGAGCAAAATCACTCC 3413
Db 3481 AGGCTGGTCCAGCAATTTATGTAAAGCTTCTAAGCCCTGGGACGAGCAAAATCACTCC 3540
QY 3414 AACAGAGCCACCTAAGTCAACAGCATTTAGCGGAGTGTGCGAGGTGATTTGGGATGACGA 3473
Db 3541 AACAGAGCCACCTAAGTCAACAGCATTTAGCGGAGTGTGCGAGGTGATTTGGGATGACGA 3600
QY 3474 CTACACCGCGAGATGCAATGAGTGGGCTTCAACAGGGCCAGATCATCAACGTCTCT 3533
Db 3601 CTACACCGCGAGATGCAATGAGTGGGCTTCAACAGGGCCAGATCATCAACGTCTCT 3660
QY 3534 CAACAGAGGAGCCCTGAGTGGTGGAAAGGAGAGTCAATGGCAAGTGGGGCTCTTCCC 3593
Db 3661 CAACAGAGGAGCCCTGAGTGGTGGAAAGGAGAGTCAATGGCAAGTGGGGCTCTTCCC 3720
QY 3594 ATCCAAATTTATGTAGTGTGACCAACAGACATGGACCCCAAGCCAGCAATG 3641
Db 3721 ATCCAAATTTATGTAGTGTGACCAACAGACATGGACCCCAAGCCAGCAATG 3768

RESULT 3

US-10-287-436A-685
; Sequence 685, Application US/10287436A
; Publication No. US20050202421A1
; GENERAL INFORMATION:
; APPLICANT: CHILDREN'S HOSPITAL MEDICAL CENTER
; TITLE OF INVENTION: METHOD FOR DIAGNOSIS AND TREATMENT OF
; TITLE OF INVENTION: RHEUMATOID ARTHRITIS
; FILE REFERENCE: 10872.514696
; CURRENT APPLICATION NUMBER: US/10/287,436A
; PRIOR FILING DATE: 2002-10-31
; PRIOR APPLICATION NUMBER: US 60/336,220
; PRIOR FILING DATE: 2001-10-31
; NUMBER OF SEQ ID NOS: 1446
; SOFTWARE: FASTSEQ for Windows Version 4.0
; SEQ ID NO 685
; LENGTH: 7247
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-287-436A-685

Query Match 63.2%; Score 3285.6; DB 9; Length 7247;

Best Local Similarity 93.8%; Pred. No. 0;

Matches 3536; Conservative 0; Mismatches 4; Indels 228; Gaps 2;

QY 102 GCGTCCCTCCACGCGCGGTGAGCGGCACTGATTTGTCCCTGGGGCGGCGAGCGGAC 161
Db 1 GCGTCCCTCCACGCGCGGTGAGCGGCACTGATTTGTCCCTGGGGCGGCGAGCGGAC 60
QY 162 CGCCCGGAGATGAGGCGTCCGATTTAGCAAGGTAAAGTAAACAGAACCATGCTCAGTTTC 221
Db 61 CGCCCGGAGATGAGGCGTCCGATTTAGCAAGGTAAAGTAAACAGAACCATGCTCAGTTTC 120
QY 222 AACACCTTTTGTGGCAGCTTGATATCTGGGCCATAAATCTGTAGAGAAAGAGCGAGCA 281
Db 121 AACACCTTTTGTGGCAGCTTGATATCTGGGCCATAAATCTGTAGAGAAAGAGCGAGCA 180
QY 282 TGNATGAGCTTCCATAGTTTAAAGCCAAATATCTGGATTCATTCTGGTGATCAAGCTAG 341
Db 181 TGNATGAGCTTCCATAGTTTAAAGCCAAATATCTGGATTCATTCTGGTGATCAAGCTAG 240
QY 342 AAACTTTTTCATCTGGGTACCTCAACCTGTTTGTAGCAGACAGATATGGGCACTAGC 401
Db 241 AAACTTTTTCATCTGGGTACCTCAACCTGTTTGTAGCAGACAGATATGGGCACTAGC 300
QY 402 TGACATGAATAATGATGGAGAAATGGATCAAGTGGAGTTTTTCCATAGCTATGAACCTAT 461
Db 301 TGACATGAATAATGATGGAGAAATGGATCAAGTGGAGTTTTTCCATAGCTATGAACCTAT 360
QY 462 CAAACTGAAGTACAGGATATCAGCTACCTCTGCACCTTCCCTCTGTCATGAAACAGCA 521
Db 361 CAAACTGAAGTACAGGATATCAGCTACCTCTGCACCTTCCCTCTGTCATGAAACAGCA 420
QY 522 ACCAGTTGCTATTCTTAGCGCACAGCATTTGGTATGGGAGGTATGCCAGCATGCCACC 581
Db 421 ACCAGTTGCTATTCTTAGCGCACCAATTTGGTATGGGAGGTATGCCAGCATGCCACC 480
QY 582 GCTTACAGCTGTTGCTCCAGTGCGCAATGGGATCCATTTCCAGTTGTTGGATGTTCTCAAC 641
Db 481 GCTTACAGCTGTTGCTCCAGTGCGCAATGGGATCCATTTCCAGTTGTTGGATGTTCTCAAC 540
QY 642 CCTAGTATCTTCTGTTCCCAACAGCATGTCGCCCTTCCCTCTGTCATGAAAGAGTTT 701
Db 541 CCTAGTATCTTCTGTTCCCAACAGCATGTCGCCCTTCCCTCTGTCATGAAAGAGTTT 600
QY 702 TATACAACTCTGCTGCAATTTGCTCATCTCTGAGGCAATTTGCCAAAGAGTTTCTCTT 761
Db 601 TATACAACTCTGCTGCAATTTGCTCATCTCTGAGGCAATTTGCCAAAGAGTTTCTCTT 660
QY 762 TAGTAGATCTGGTCCAGGCTCAAACTAAACATTAACAAAGGCAAGTCAATTTGA 821

Db 661 TAGTAGATCTGCTCCAGGGTCAAACTAAACATAAAATTAACAAAGGCAAGTCAATTGA 720
Qy 822 TGTGGCCAGTGTCCACCAAGTGGCAGAGTGGGCTGTTCTCAGTCATCAAGACTGAAATA 881
Db 721 TGTGGCCAGTGTCCACCAAGTGGCAGAGTGGGCTGTTCTCAGTCATCAAGGCTGAATA 780
Qy 882 CAGGCAATTAATCAATAGTCATGACAAACTATAGAGTGACACTTAAACAGGTCCCAAGC 941
Db 781 CAGGCAATTAATCAATAGTCATGACAAACTATAGAGTGACACTTAAACAGGTCCCAAGC 840
Qy 942 RAGAACTATTCTTATGTCAGTCAAGTTTACCACAGGCTCAGCTGGCTTCAATATGAATCT 1001
Db 841 AAGAACTATTCTTATGTCAGTCAAGTTTACCACAGGCTCAGCTGGCTTCAATATGAATCT 900
Qy 1002 TTCTGACATTTGATCAAGATGGAACCTTACAGCAGAGGAATTTATCTGGCAATGCACCT 1061
Db 901 TTCTGACATTTGATCAAGATGGAACCTTACAGCAGAGGAATTTATCTGGCAATGCACCT 960
Qy 1062 CATTGATGTAGTATGTCTGGCCAACTGCACTGCCACTGCTGCTGCTCCAGAAATACATTC 1121
Db 961 CATTGATGTAGTATGTCTGGCCAACTGCACTGCCACTGCTGCTGCTCCAGAAATACATTC 1020
Qy 1122 ACCTTCTTTTAGAAGTTCGATCTGGCAGTGGTATATCTGTCAATAGCTCAACATCTGT 1181
Db 1021 ACCTTCTTTTAGAAGTTCGATCTGGCAGTGGTATATCTGTCAATAGCTCAACATCTGT 1080
Qy 1182 AGATCAGAGGCTACACAGAGAACCCAGTTTGTAGAAGATGAACAACTAAATTAGAAAGAA 1241
Db 1081 AGATCAGAGGCTACACAGAGAACCCAGTTTGTAGAAGATGAACAACTAAATTAGAAAGAA 1140
Qy 1242 ATTACTGTAAAGTTTGAAGATAAGNAGCGGAGAACTTTGAACTGGGCAACTGGAACCT 1301
Db 1141 ATTACTGTAAAGTTTGAAGATAAGNAGCGGAGAACTTTGAACTGGGCAACTGGAACCT 1200
Qy 1302 GGAGAAACGAAGCAAGCTCTCTGGAAACAGCAGCGCAAGGAGCAGAGCGCTGGCCCA 1361
Db 1201 GGAGAAACGAAGCAAGCTCTCTGGAAACAGCAGCGCAAGGAGCAGAGCGCTGGCCCA 1260
Qy 1362 GCTGGAGCGGGCGGAGCAGAGAGGAGGAGCGGTGAGCGCCAGGAGCAAGAGCGCAAAAG 1421
Db 1261 GCTGGAGCGGGCGGAGCAGAGAGGAGGAGCGGTGAGCGCCAGGAGCAAGAGCGCAAAAG 1320
Qy 1422 ACAACTGGAACTGGAGAACCACTGGAAAAGCAGCGGAGCTAGAAACGCGCAGAGAGGA 1481
Db 1321 ACAACTGGAACTGGAGAACCACTGGAAAAGCAGCGGAGCTAGAAACGCGCAGAGAGGA 1380
Qy 1482 GGAGAGGAGGAAAGAAATTTGAGAGCGGAGAGGCTGCAAAACGGAACTTTGAAAGGCAAG 1541
Db 1381 GGAGAGGAGGAAAGAAATTTGAGAGCGGAGAGGCTGCAAAACGGAACTTTGAAAGGCAAG 1440
Qy 1542 ACAACTGGAGTGGGAAACCGAATCGAAGGCAAGAACTTAAATCAAAAGAAAACAAAGAAC 1601
Db 1441 ACAACTGGAGTGGGAAACCGAATCGAAGGCAAGAACTTAAATCAAAAGAAAACAAAGAAC 1500
Qy 1602 AGAGGACATAGTTTACTGAAAGCAAGAAAGAACTTTGGAAATTTGAAATTTAGAGCTCT 1661
Db 1501 AGAGGACATAGTTTACTGAAAGCAAGAAAGAACTTTGGAAATTTGAAATTTAGAGCTCT 1560
Qy 1662 AAATGATAAAAGCACTCAATAGAGGAACTTCAAGATATCAGATGTGCAATTTGACAC 1721
Db 1561 AAATGATAAAAGCACTCAATAGAGGAACTTCAAGATATCAGATGTGCAATTTGACAC 1620
Qy 1722 CCAAGGCAAGAAATTTAGAGCAGCAAAACAAATCTAGAGAGTTGAGAAATTTGCCAAATCAC 1781
Db 1621 CCAAGGCAAGAAATTTAGAGCAGCAAAACAAATCTAGAGAGTTGAGAAATTTGCCAAATCAC 1680
Qy 1782 CCATCTAGACCAACAAATTTACAGGAATCTCAGCAAAATGCTTGGAGACTTTATTCAGAA 1841
Db 1681 CCATCTAGACCAACAAATTTACAGGAATCTCAGCAAAATGCTTGGAGACTTTATTCAGAA 1740
Qy 1842 ACAGATACTCAATGACCAATTAACCAAGTTTACAGAGAAACAGTTTGCACAGAGATTCAC 1901
Db 1741 ACAGATACTCAATGACCAATTAACCAAGTTTACAGAGAAACAGTTTGCACAGAGATTCAC 1800

Qy 1902 TGTACACTTAAAGAGCCTTTAGAGCAAAAGAACTAGCTCGGAGCACTTACGAGACCA 1961
Db 1801 TGTACACTTAAAGAGCCTTTAGAGCAAAAGAACTAGCTCGGAGCACTTACGAGACCA 1860
Qy 1962 ACTGGATGAAGTGGAGAAAGAACTAGATCAAAACTACAGGAGATTGATATTTTCAATAA 2021
Db 1861 ACTGGATGAAGTGGAGAAAGAACTAGATCAAAACTACAGGAGATTGATATTTTCAATAA 1920
Qy 2022 TCAGCTGAAGAACTTAAGAGAAATACAACTAAGCAACAACTCCAGAAAGCAAAAGTCCAT 2081
Db 1921 TCAGCTGAAGAACTTAAGAGAAATACAACTAAGCAACAACTCCAGAAAGCAAAAGTCCAT 1980
Qy 2082 GGAGGCTGAACGACTGAAGAAACAGAAAGAAACAGAAAGATCATAGAAATTAGAAAAACA 2141
Db 1981 GGAGGCTGAACGACTGAAGAAACAGAAAGAAACAGAAAGATCATAGAAATTAGAAAAACA 2040
Qy 2142 AAAAGAAAGCCCAAGACGAGCTCAGGAAAGGACAAAGCAGTGGCTGGAGCATGTGCA 2201
Db 2041 AAAAGAAAGCCCAAGACGAGCTCAGGAAAGGACAAAGCAGTGGCTGGAGCATGTGCA 2100
Qy 2202 GCAGAGGACGAGCATCAGAGACCAAGAAAACTCCACGAAGAGGAAAACTGAAAAAGGA 2261
Db 2101 GCAGAGGACGAGCATCAGAGACCAAGAAAACTCCACGAAGAGGAAAACTGAAAAAGGA 2160
Qy 2262 GGAGAGTGTCAAAAGAAAGGATGGCGAGGAAAAAGGCAAAACAGGAGCACAAGACAAGCT 2321
Db 2161 GGAGAGTGTCAAAAGAAAGGATGGCGAGGAAAAAGGCAAAACAGGAGCACAAGACAAGCT 2220
Qy 2322 GGGTCGGCTTTTCCATCAACCAAGAACCAAGCAGCTGTAAGCAGCTGTCAGGCAACCTGGTC 2381
Db 2221 GGGTCGGCTTTTCCATCAACCAAGAACCAAGCAGCTGTAAGCAGCTGTCAGGCAACCTGGTC 2280
Qy 2382 CACTGCAGAAAAAGGTCCACTTTACCATTTCTGCAAGGAAAAATGTAAAAAGTGGTGTATTA 2441
Db 2281 CACTGCAGAAAAAGGTCCACTTTACCATTTCTGCAAGGAAAAATGTAAAAAGTGGTGTATTA 2340
Qy 2442 CCGGCACTGTACCCCTTTGAATCCAGAAAGCATGATCAATCATATCCAGCCAGGAGA 2501
Db 2341 CCGGCACTGTACCCCTTTGAATCCAGAAAGCATGATCAATCATATCCAGCCAGGAGA 2400
Qy 2502 CATAGTCAT-----GGTGGATGAAAGCCAAACTGGAGAAACCGCGCTGGCT 2546
Db 2401 CATAGTCATGGTTAAAGGGGAATGGTGGATGAAAGCCAACTGGAGAAACCGCGCTGGCT 2460
Qy 2547 TGGAGGAGAAATTAAGAGGAAAGACAGGGTGGTTCCTGCAAACTATGAGAGAAAAATCCC 2606
Db 2461 TGGAGGAGAAATTAAGAGGAAAGACAGGGTGGTTCCTGCAAACTATGAGAGAAAAATCCC 2520
Qy 2607 AGAAAAAGAGTTCCCGCTCCAGTGAACCAAGTGAATGATTCACATCTGCCCTGCCCC 2666
Db 2521 AGAAAAAGAGTTCCCGCTCCAGTGAACCAAGTGAATGATTCACATCTGCCCTGCCCC 2580
Qy 2667 CAAACTGGCTTGGTGAAGACCCCGCTTTGGCAGTAACTCTTCAGAGCCCTCCAC 2726
Db 2581 CAAACTGGCTTGGTGAAGACCCCGCTTTGGCAGTAACTCTTCAGAGCCCTCCAC 2640
Qy 2727 GACCCCTAAATACTGGGCGGACTTCAGCTCAAGTGGGCGCCACAGCAAGATGAGAAACC 2786
Db 2641 GACCCCTAAATACTGGGCGGACTTCAGCTCAAGTGGGCGCCACAGCAAGATGAGAAACC 2700
Qy 2787 AGAAACGGAATACTGGGATGATGGGAGCCCGCTCTCTCACCGTTCCAAGTGGCGG 2846
Db 2701 AGAAACGGAATACTGGGATGATGGGAGCCCGCTCTCTCACCGTTCCAAGTGGCGG 2760
Qy 2847 CCAGTTAAGGAGAGGTCGCTTTTACTCCAGCCACGGCCACTGGCTCTCCCGCTCTCC 2906
Db 2761 CCAGTTAAGGAGAGGTCGCTTTTACTCCAGCCACGGCCACTGGCTCTCCCGCTCTCC 2820
Qy 2907 TGTGTCAGGCCAGGGTGAAGAGTGGAGGGGTACAAGCTCAAGCCCTATATCTTCTTGGAG 2966
Db 2821 TGTGTCAGGCCAGGGTGAAGAGTGGAGGGGTACAAGCTCAAGCCCTATATCTTCTTGGAG 2880

| | | | | | | | |
|----|------|--|------|----|------|---|------|
| QY | 794 | CTAAATTACAAAGGCACAGTCAATTTGATGTGGCCAGTCTCCACC - AGTGCAGAGTGG | 852 | QY | 1858 | CAATTAAAAAAGTTTCAGCAGAAACAGTTTGCACAGAGATTCACTTTGTTACACTTAAAAAG - | 1916 |
| Db | 850 | CTAAATTACAAAGGCACAGTCAATTTGATGTGGCCAGTCTCCACCAGTGGCAGAGTGG | 909 | Db | 1930 | CAATTAAAAAAGTTTCAGCAGAAACAGTTTGCACAGAGATTCACTTTGTTACACTTAAAAAG | 1989 |
| QY | 853 | GCTGTTCTCAGTCATCAGACTGAATACAAA - GGCATTAATCAATAGTCATGACAAA | 909 | QY | 1917 | AGCCTTTAGAAGCAAAAGAACTAGCTCGGCAGCACCTACGAGACCACTTGGATGGAAGTGA | 1976 |
| Db | 910 | GCTGTTCTCAGTCATCAGACTGAATACAAA - GGCATTAATCAATAGTCATGACAAA | 969 | Db | 1990 | AGCCTTTAGAAGCAAAAGAACTAGCTCGGCAGCACCTACGAGACCACTTGGATGGAAGTGA | 2049 |
| QY | 910 | AC - TATGAGTGACACTT - AACAGGTCCCAAGCAAGAACTATTCTTATGTCAGTCAAG | 965 | QY | 1977 | GAAGAAATCTAGATCAAAACTACAGGAGATTGATATTTTCAATAATAGCTGAAGAACT | 2036 |
| Db | 970 | ACCTAATGAGTGACACTTTTAAAGGTCCCAAGCAAGAACTATTCTTATGTCAGTCAAG | 1029 | Db | 2050 | GAAGAAATCTAGATCAAAACTACAGGAGATTGATATTTTCAATAATAGCTGAAGAACT | 2109 |
| QY | 966 | TTTACCACAGGCTCAGCTGGCTTCAATATGGAATCTTTCTGATGATCAAGATGGA | 1025 | QY | 2037 | RAGAGAAATACAAATAGCAACACTCCAGNAGCAAAAGTCCATGGAGGCTGAACGACT | 2096 |
| Db | 1030 | TTTACCACAGGCTCAGCTGGCTTCAATATGGAATCTTTCTGATGATCAAGATGGA | 1089 | Db | 2110 | RAGAGAAATACAAATAGCAACACTCCAGNAGCAAAAGTCCATGGAGGCTGAACGACT | 2169 |
| QY | 1026 | ACTTACAGCAGAGGAATTTATCTTGGCAATGCACCTCATTTGATAGCTATGCTGGCCA | 1085 | QY | 2097 | GAACACAGAAAGAAACAAAGAACGAAAGATCATAGAAATTTAGAAAAACAAAAGAAAGAACGCCA | 2156 |
| Db | 1090 | ACTTACAGCAGAGGAATTTATCTTGGCAATGCACCTCATTTGATAGCTATGCTGGCCA | 1149 | Db | 2170 | GAACACAGAAAGAAACAAAGAACGAAAGATCATAGAAATTTAGAAAAACAAAAGAAAGAACGCCA | 2229 |
| QY | 1086 | ACCACCTGCCACCTGCTCCCTCCAGAAATACATTCACACCTTCTTTTAGAAGAGTTTGATC | 1145 | QY | 2157 | AAGACGAGCTCAGGAAAGGGAACAGCAGTGGCTGGAGCATGTGTCAGCAGGAGGACGAGCA | 2216 |
| Db | 1150 | ACCACCTGCCACCTGCTCCCTCCAGAAATACATTCACACCTTCTTTTAGAAGAGTTTGATC | 1209 | Db | 2230 | AAGACGAGCTCAGGAAAGGGAACAGCAGTGGCTGGAGCATGTGTCAGCAGGAGGACGAGCA | 2289 |
| QY | 1146 | TGGCAGTGGTATATCTGTATAGCTCAACATCTGTGATATCAGAGGCTACAGAGAAAC | 1205 | QY | 2217 | TCAGAGACCAAGAAACCTCCAGNAGAGGAAACCTGAAAGGGAGGAGAGTGTCAAAA | 2276 |
| Db | 1210 | TGGCAGTGGTATATCTGTATAGCTCAACATCTGTGATATCAGAGGCTACAGAGAAAC | 1269 | Db | 2290 | TCAGAGACCAAGAAACCTCCAGNAGAGGAAACCTGAAAGGGAGGAGAGTGTCAAAA | 2349 |
| QY | 1206 | AGTTTTAGAGATGAACAAACAACTAGAAAAGAAATTTACCTGTAAACGTTTGAAGATAA | 1265 | QY | 2277 | GAAGGATGGC - GAGGAAAAAGGCAACAGGAAGCAACAGACAAGCTGGGTTCGGCTTTTCC | 2335 |
| Db | 1270 | AGTTTTAGAGATGAACAAACAACTAGAAAAGAAATTTACCTGTAAACGTTTGAAGATAA | 1329 | Db | 2350 | GAAGGATGGCAGAGGAAAGGCAACAGGAAGCAACAGACAAGCTGGGTTCGGCTTTTCC | 2409 |
| QY | 1266 | GAAGCGGGAAGACTTTGAAACGTGGCAACCTGGAACCTGGAGAAACGAAGGCAAGCTCTCCT | 1325 | QY | 2336 | ATCAACCAAGAAACAGCTAAGCAGTGTTCAGGCAACCTTGGTCTCACTGCAGAAAAAG | 2395 |
| Db | 1330 | GAAGCGGGAAGACTTTGAAACGTGGCAACCTGGAACCTGGAGAAACGAAGGCAAGCTCTCCT | 1389 | Db | 2410 | ATCAACCAAGAAACAGCTAAGCAGTGTTCAGGCAACCTTGGTCTCACTGCAGAAAAAG | 2469 |
| QY | 1326 | GAAGCAGCAGCAGAGGAGAGAGCGCTTGGCCAGCTGGAGCGGGCGGAGCAGAGAG | 1385 | QY | 2396 | GTCCACTTTACCAATTTCTGCACAGGAAATGTAAAGTGTGTATTACCGGGCACTGTACC | 2455 |
| Db | 1390 | GAAGCAGCAGCAGAGGAGAGAGCGCTTGGCCAGCTGGAGCGGGCGGAGCAGAGAG | 1449 | Db | 2470 | GTCCACTTTACCAATTTCTGCACAGGAAATGTAAAGTGTGTATTACCGGGCACTGTACC | 2529 |
| QY | 1386 | GAAGGAGCTGAGCGCCAGGAGCAGAGCGCAAGAGCAACCTGGAACCTGGAGAGCAACT | 1445 | QY | 2456 | CTTTTGAATTCAGAAAGCCATGATGAAATCACTATCCAGCAGGAGACATAGTCAT - | 2510 |
| Db | 1450 | GAAGGAGCTGAGCGCCAGGAGCAGAGCGCAAGAGCAACCTGGAACCTGGAGAGCAACT | 1509 | Db | 2530 | CTTTTGAATTCAGAAAGCCATGATGAAATCACTATCCAGCAGGAGACATAGTCATGGTTA | 2589 |
| QY | 1446 | GGAAAGCAGCGGAGCTAGAAACGCGCAGAGAGAGGAGAGAGGAGGAGAAATTCAGAG | 1505 | QY | 2511 | -----GGTGGATGAAGCCAAAACCTGGAGAAACCGCGCTGGCTTGGAGAGAGATTAA | 2560 |
| Db | 1510 | GGAAAGCAGCGGAGCTAGAAACGCGCAGAGAGAGGAGAGGAGGAGGAGGAGAAATTCAGAG | 1569 | Db | 2590 | GAAGGGAATTTGGTGGATGAAGCCAAAACCTGGAGAAACCGCGCTGGCTTGGAGAGAGATTAA | 2649 |
| QY | 1506 | GCGAGAGGCTGCAAAACGGGAAC - TGAAGGCAACGACAACTTGTGAGTGGGAACGGAATC | 1564 | QY | 2561 | RAGGAAAGCAGGCTGGTTCCCTGCAAACTATGCAGAGAAAAATCCAGAAAAATGAGGTTTC | 2620 |
| Db | 1570 | GCGAGAGGCTGCAAAACGGGAAC - TGAAGGCAACGACAACTTGTGAGTGGGAACGGAATC | 1629 | Db | 2650 | RAGGAAAGCAGGCTGGTTCCCTGCAAACTATGCAGAGAAAAATCCAGAAAAATGAGGTTTC | 2709 |
| QY | 1565 | GAAGGCAAGAACTTCTAAATCAAGAAACAAAGAACAGAGGACATAGTTGTACTGAAG | 1624 | QY | 2621 | CCGCTCCAGTGAAACAGTGACTGATTCAACATCTGCCCTGCCCTGCCCTGCCCTGCCCTGCC | 2680 |
| Db | 1630 | GAGAGTCCAACTACTAAATCAAGAAACAAAGAACAGAGGACATAGTTGTACTGAAG | 1689 | Db | 2710 | CCGCTCCAGTGAAACAGTGACTGATTCAACATCTGCCCTGCCCTGCCCTGCCCTGCCCTGCC | 2769 |
| QY | 1625 | CAAGAAAAAGACTTTTGAATTTGAATTTGAAGCTCTAAATGATAAAAAAGCATCAACTAG | 1684 | QY | 2681 | GTGAGACCCCGCCCTTTTGGCAGTAACTCTTTTGAAGCCCTCCAGCCCTCCAGCCCTTAATACT | 2740 |
| Db | 1690 | CAAGAAAAAGACTTTTGAATTTGAATTTGAAGCTCTAAATGATAAAAAAGCATCAACTAG | 1749 | Db | 2770 | GTGAGACCCCGCCCTTTTGGCAGTAACTCTCTCACCGTTTCAAGTGCAGGCGCAGTTAAGGCA | 2829 |
| QY | 1685 | AAGGGAACCTCAAGATATCAGATGTCGATGACCAACCAAGGCAAGAAATTTGAGAGCA | 1744 | QY | 2741 | GGGCGGACTTCAGCTCCAGTGGCCACCCAGCAGCAAGATGAGAAAAACGAGAAAAACGATAACT | 2800 |
| Db | 1750 | AAGGGAACCTCAAGATATCAGATGTCGATGACCAACCAAGGCAAGAAATTTGAGAGCA | 1809 | Db | 2830 | GGGCGGACTTCAGCTCCAGTGGCCACCCAGCAGCAAGATGAGAAAAACGAGAAAAACGATAACT | 2889 |
| QY | 1745 | CAAGAAAACTAGAGAGTTGAGATTTGCCG - AAATCACCCATCTACAGCAAACTT - A | 1800 | QY | 2801 | GGGATGATGGGAGCCAGCCCTTCTCTCACCGTTTCAAGTGCAGGCGCAGTTAAGGCA | 2859 |
| Db | 1810 | CAAGAAAACTAGAGAGTTGAGATTTGCCG - AAATCACCCATCTACAGCAAACTT - A | 1869 | Db | 2890 | GGGATGATGGGAGCCAGCCCTTCTCTCACCGTTTCAAGTGCAGGCGCAGTTAAGGCA | 2949 |
| QY | 1801 | CAGGAATCTCAGCAAA - --TGCTTGAAGACTTTATTCAGAAAAACAGATCTCAATGAC | 1857 | QY | 2860 | AGGTGCGCTTTTACTCCAGCCAGGCGCACTGCTCTCCCGCTCTCTGCTGCTAGGCCAG | 2919 |
| Db | 1870 | ATGGAATCTCAGCAAAATGCTTTTGAAGAACTTTATTCAGAAAAACAGATCTCAATGAC | 1929 | Db | 2950 | AGGTGCGCTTTTACTCCAGCCAGGCGCACTGCTCTCCCGCTCTCTGCTGCTAGGCCAG | 3009 |

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|----|------|---|------|----|------|---|------|
| QY | 795 | TAAATTTACAAAAGGCACAGTCATTTGATGTGGCCAGTGTCCCAACAGTGGCAGAGTGGGC | 854 | QY | 1863 | AAAAAAGCTTCAGCAGAACAGCTTTGCACAGAGATTCACCTGTTACACTTAAAGAGCCCTT | 1922 |
| DB | 864 | TAAATTTACAAAAGGCACAGTCATTTGATGTGGCCAGTGTCCCAACAGTGGCAGAGTGGGC | 923 | DB | 1944 | AAAAAAGCTTCAGCAGAACAGCTTTGCACAGAGATTCACCTGTTACACTTAAAGAGCCCTT | 2003 |
| QY | 855 | TGTTCTCAGTCATCAAGACTGAAATACAGGCAATTTATTCATAGTCATGACAAACTAT | 914 | QY | 1923 | AGAAAGAAAAGAACTTAGCTCGGCAGCCTACGAGACCAACTGGATGAAGTGGAGAAAAG | 1982 |
| DB | 924 | TGTTCTCAGTCATCAAGACTGAAATACAGGCAATTTATTCATAGTCATGACAAACTAT | 983 | DB | 2004 | AGAAAGAAAAGAACTTAGCTCGGCAGCCTACGAGACCAACTGGATGAAGTGGAGAAAAG | 2063 |
| QY | 915 | GAGTGGACACTTAAC-----AGGTCCCCAAGCAAGAACTATTCTTATGCGAGTC | 962 | QY | 1983 | AACATAGTCAAAACTACAGGAGATTGATTTTTCAATATCAGCTGAAGGAACCTAAGAGA | 2042 |
| DB | 984 | GAGTGGACACTTAACAGGTTCTGTTAGGTCCCCAAGCAAGAACTATTCTTATGCGAGTC | 1043 | DB | 2064 | AACATAGTCAAAACTACAGGAGATTGATTTTTCAATATCAGCTGAAGGAACCTAAGAGA | 2123 |
| QY | 963 | AAAGTTTACACAGGCTCAGCTGGCTCAATATGGAATCTTTCTGACATTTGATCAAGATGG | 1022 | QY | 2043 | AATACACATATAGCAACCAACTCCGAAGCAAAAGTCCATGAGGCTGAACGACTGAACA | 2102 |
| DB | 1044 | AAAGTTTACACAGGCTCAGCTGGCTCAATATGGAATCTTTCTGACATTTGATCAAGATGG | 1103 | DB | 2124 | AATACACATATAGCAACCAACTCCGAAGCAAAAGTCCATGAGGCTGAACGACTGAACA | 2183 |
| QY | 1023 | AAAACTTACAGCAGAGGAATTTATCTGCGCAATGCACTCATTTGATGATGCTGCTGG | 1082 | QY | 2103 | GAAAGAACAAAGAACGAAAGATCATAGAAATTAGAAAAACAAAAGAAAGGCCCAAAGAGC | 2162 |
| DB | 1104 | AAAACTTACAGCAGAGGAATTTATCTGCGCAATGCACTCATTTGATGATGCTGCTGG | 1163 | DB | 2184 | GAAAGAACAAAGAACGAAAGATCATAGAAATTAGAAAAACAAAAGAAAGGCCCAAAGAGC | 2243 |
| QY | 1083 | CCAAACCACTGCCACCTGCTCTGCTCCAGAATACATTCACACCTCTTTTTAGAGAGTTCCG | 1142 | QY | 2223 | ACCAAGAAAACCTCCACGAAGAGGAAAACTGAAAAGGAGGAGAGTGTCAAAAAGAAAGGA | 2282 |
| DB | 1164 | CCAAACCACTGCCACCTGCTCTGCTCCAGAATACATTCACACCTCTTTTTAGAGAGTTCCG | 1223 | DB | 2304 | ACCAAGAAAACCTCCACGAAGAGGAAAACTGAAAAGGAGGAGAGTGTCAAAAAGAAAGGA | 2363 |
| QY | 1143 | ATCTGCGAGTGTATATCTGTGTAATAGCTCAACATCTGTAGATCAGAGGCTACCGAGGA | 1202 | QY | 2283 | TGGCAGGAAAAGGCAACAGGAAGCAAGAACAGCAAGCTGGGTCCGCTTTTCCATCAACA | 2342 |
| DB | 1224 | ATCTGCGAGTGTATATCTGTGTAATAGCTCAACATCTGTAGATCAGAGGCTACCGAGGA | 1283 | DB | 2364 | TGGCAGGAAAAGGCAACAGGAAGCAAGAACAGCAAGCTGGGTCCGCTTTTCCATCAACA | 2423 |
| QY | 1203 | ACCAGTTTTAGAGATGAAACAACTATTAGAAAAGAAATACCTGTAACGTTTGAAGA | 1262 | QY | 2403 | TACCAATTTCTGCACAGGAAAATGTAAGTGTGTATTACCGGGCACTGTACCCCTTTGA | 2462 |
| DB | 1284 | ACCAGTTTTAGAGATGAAACAACTATTAGAAAAGAAATACCTGTAACGTTTGAAGA | 1343 | DB | 2484 | TACCAATTTCTGCACAGGAAAATGTAAGTGTGTATTACCGGGCACTGTACCCCTTTGA | 2543 |
| QY | 1263 | TAAAGACCGGGAGAACTTTGAACGTGGCAACCTGGAACTGGAGAAACGAAGGCAAGCTCT | 1322 | QY | 2463 | ATCCAGAAAGCCATGATGAATCACTATCCAGCAGGAGACATAGTCATGGTGGATGAAG | 2522 |
| DB | 1344 | TAAAGACCGGGAGAACTTTGAACGTGGCAACCTGGAACTGGAGAAACGAAGGCAAGCTCT | 1403 | DB | 2544 | ATCCAGAAAGCCATGATGAATCACTATCCAGCAGGAGACATAGTCATGGTGGATGAAG | 2603 |
| QY | 1323 | CCTGGAACAGCAGCGCAAGGAGGAGGAGCGCTTGGCCCACTGGAGCGGGCGGAGGAGGA | 1382 | QY | 2523 | CCAAACTGGAGAACCCCGCTGGCTTGGAGGAGAAATTAAGAAAGAAAGACAGGCTGGTCCC | 2582 |
| DB | 1404 | CCTGGAACAGCAGCGCAAGGAGGAGGAGGCGCTTGGCCCACTGGAGCGGGCGGAGGAGGA | 1463 | DB | 2604 | CCAAACTGGAGAACCCCGCTGGCTTGGAGGAGAAATTAAGAAAGAAAGACAGGCTGGTCCC | 2663 |
| QY | 1383 | GAGGAAGAGCGTGTAGCGCCAGAGCAAGAGCGCAAAAGACAACTGGAACTGGAGAGCA | 1442 | QY | 2583 | TGCAAACTATGCAGAGAAAATCCCAAGAAAATGAGGTTCCCGCTCCAGTGAACCAAGTGAC | 2642 |
| DB | 1464 | GAGGAAGAGCGTGTAGCGCCAGAGCAAGAGCGCAAAAGACAACTGGAACTGGAGAGCA | 1523 | DB | 2664 | TGCAAACTATGCAGAGAAAATCCCAAGAAAATGAGGTTCCCGCTCCAGTGAACCAAGTGAC | 2723 |
| QY | 1443 | ACTGGAAGAGCAGCGGAGCTAGAACGGCAGAGAGAGGAGGAGGAGGAGGAGAAATTTGA | 1502 | QY | 2643 | TGATTTCAACATCTGCCCTTGGCCCAAACTGGCTTGGCTGAGAGCCCGCCCTTTTGGC | 2702 |
| DB | 1524 | ACTGGAAGAGCAGCGGAGCTAGAACGGCAGAGAGAGGAGGAGGAGGAGGAGGAGAAATTTGA | 1583 | DB | 2724 | TGATTTCAACATCTGCCCTTGGCCCAAACTGGCTTGGCTGAGAGCCCGCCCTTTTGGC | 2783 |
| QY | 1503 | GAGGCGAGAGGCTGCAAAAAGGGAACTTGAAGGCAACGACAACTTGAAGTGGGAACGGAA | 1562 | QY | 2703 | AGTAACCTTTTCAGAGCCCTCCACGACCCCTTAATTAACCTGGGCCGACTTTCAGCTCCACGCTG | 2762 |
| DB | 1584 | GAGGCGAGAGGCTGCAAAAAGGGAACTTGAAGGCAACGACAACTTGAAGTGGGAACGGAA | 1643 | DB | 2784 | AGTAACCTTTTCAGAGCCCTCCACGACCCCTTAATTAACCTGGGCCGACTTTCAGCTCCACGCTG | 2843 |
| QY | 1563 | TCGAAGGCAAGAACTACTAAATCAAGAAACAAAGAAACAAAGGACATAGTTGACTGAA | 1622 | QY | 2763 | GCCCAACAGCAGCAATGAGAACCAAGAAACGGAATTAACCTGGGATGATGGGAGCCAGCC | 2822 |
| DB | 1644 | TCGAAGGCAAGAACTACTAAATCAAGAAACAAAGAAACAAAGGACATAGTTGACTGAA | 1703 | DB | 2844 | GCCCAACAGCAGCAATGAGAACCAAGAAACGGAATTAACCTGGGATGATGGGAGCCAGCC | 2903 |
| QY | 1623 | AGCAAGAAAGAAAGACTTTTGGAAATTTGAATTTAGAGCTCTAAATGATPAAAAAGCATCACT | 1682 | QY | 2823 | CTCTCTCACCGTTTCAAGTGGCGGCAAGTTAAGGAGAGGTCGCTTTTACTCCAGCCAC | 2882 |
| DB | 1704 | AGCAAGAAAGAAAGACTTTTGGAAATTTGAATTTAGAGCTCTAAATGATPAAAAAGCATCACT | 1763 | DB | 2904 | CTCTCTCACCGTTTCAAGTGGCGGCAAGTTAAGGAGAGGTCGCTTTTACTCCAGCCAC | 2963 |
| QY | 1683 | AGAGGGAACCTTCAAGATATCAGATGTGATTTGACCAACCAAGGCAAGAAATTTGAGAG | 1742 | QY | 2883 | GGCCTAGCTCTCCCGTCTCCTGTGCTAGGCCAGGGTGA | 2924 |
| DB | 1764 | AGAGGGAACCTTCAAGATATCAGATGTGATTTGACCAACCAAGGCAAGAAATTTGAGAG | 1823 | DB | 2964 | GGCCTAGCTCTCCCGTCTCCTGTGCTAGGCCAGGGTGA | 3005 |
| QY | 1743 | CACAAACAAATCTAGAGAGTTGAGAAATTCGGAATTCACCCATCTACAGCAACAAATTACA | 1802 | | | | |
| DB | 1824 | CACAAACAAATCTAGAGAGTTGAGAAATTCGGAATTCACCCATCTACAGCAACAAATTACA | 1883 | | | | |
| QY | 1803 | GGAACTCTAGCAAAATGCTTTGGAAGACTTTATTCAGAAAAACAGATACTCAATGACCAATT | 1862 | | | | |
| DB | 1884 | GGAACTCTAGCAAAATGCTTTGGAAGACTTTATTCAGAAAAACAGATACTCAATGACCAATT | 1943 | | | | |

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| US-09-764-875-88 | | | | | | | | | |
| ; Sequence 88, Application US/09764875 | | | | | | | | | |
| ; Publication No. US20040018969A1 | | | | | | | | | |
| ; GENERAL INFORMATION: | | | | | | | | | |
| ; APPLICANT: Rosen et al. | | | | | | | | | |
| ; TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies | | | | | | | | | |
| ; FILE REFERENCE: PJ202 | | | | | | | | | |
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| ; CURRENT FILING DATE: 2001-01-17 | | | | | | | | | |
| ; Prior application data removed - consult PALM or file wrapper | | | | | | | | | |
| ; NUMBER OF SEQ ID NOS: 1249 | | | | | | | | | |
| ; SOFTWARE: PatentIn Ver. 2.0 | | | | | | | | | |
| ; SEQ ID NO 88 | | | | | | | | | |
| ; LENGTH: 3319 | | | | | | | | | |
| ; TYPE: DNA | | | | | | | | | |
| ; ORGANISM: Homo sapiens | | | | | | | | | |
| US-09-764-875-88 | | | | | | | | | |
| Query Match 54.7%; Score 2843.2; DB 3; Length 3319; | | | | | | | | | |
| Best Local Similarity 99.5%; Pred. No. 0; | | | | | | | | | |
| Matches 2867; Conservative 0; Mismatches 3; Indels 12; Gaps 1; | | | | | | | | | |
| QY | 55 | GACGGACAGAGGCGGGGGGATGTTGCGGGGCTCGGGCTCTCGGCTCCCTCCCGAG | 114 | | | | | | |
| DB | 6 | GAGGGACAGAGGCGGGGGGATGTTGCGGGGCTCGGGCTCTCGGCTCCCTCCCGAG | 65 | | | | | | |
| QY | 115 | CGGCGGCTGAGCGGCACTGATTTGTCTCCCTGGGCGGACCGCGGACCCCGCGAGATGA | 174 | | | | | | |
| DB | 66 | CGGCGGCTGAGCGGCACTGATTTGTCTCCCTGGGCGGACCGCGGACCCCGCGAGATGA | 125 | | | | | | |
| QY | 175 | GGCGTGAATTAGCAAGGTAAGTAACAGAACCATGGCTCAGTTTCCACACCTTTTGGT | 234 | | | | | | |
| DB | 126 | GGCGTGAATTAGCAAGGTAAGTAACAGAACCATGGCTCAGTTTCCACACCTTTTGGT | 185 | | | | | | |
| QY | 235 | GGCAGCTGATATCTGGGCGAATACTGTAGAGAAAGAGCGAGCATGATCAGCAGTTTC | 294 | | | | | | |
| DB | 186 | GGCAGCTGATATCTGGGCGAATACTGTAGAGAAAGAGCGAGCATGATCAGCAGTTTC | 245 | | | | | | |
| QY | 295 | CATAGTTTAAAGCAATATCTGGATTCATTACTGTTGATCAAGCTAGAACTTTTTTTTT | 354 | | | | | | |
| DB | 246 | CATAGTTTAAAGCAATATCTGGATTCATTACTGTTGATCAAGCTAGAACTTTTTTTTT | 305 | | | | | | |
| QY | 355 | CAATCTGGGTTACTCAACTCTGTTTGTAGCAAGATATGGCACTAGCTGATGAATAAT | 414 | | | | | | |
| DB | 306 | CAATCTGGGTTACTCAACTCTGTTTGTAGCAAGATATGGCACTAGCTGATGAATAAT | 365 | | | | | | |
| QY | 415 | GATGGAGATGATCAAGTGGAGTTTCCATAGCTATGAACTTATCAAACTGAAGCTA | 474 | | | | | | |
| DB | 366 | GATGGAGATGATCAAGTGGAGTTTCCATAGCTATGAACTTATCAAACTGAAGCTA | 425 | | | | | | |
| QY | 475 | CAAGGATATCAGCTACCTCTGCACCTTCCCTCTGTCTATGAAACAGCAACAGTTGCTATT | 534 | | | | | | |
| DB | 426 | CAAGGATATCAGCTACCTCTGCACCTTCCCTCTGTCTATGAAACAGCAACAGTTGCTATT | 485 | | | | | | |
| QY | 535 | TCATGCCACAGCATTTGGTATGGGAGGTATGCCAGCATGCCACCGCTTACAGCTGTT | 594 | | | | | | |
| DB | 486 | TCATGCCACAGCATTTGGTATGGGAGGTATGCCAGCATGCCACCGCTTACAGCTGTT | 545 | | | | | | |
| QY | 595 | GCTCCAGTCCAAATGGGATCCATTCCAGTTGTTGGAAATGTTCCCAACCTAGTATCTTCT | 654 | | | | | | |
| DB | 546 | GCTCCAGTCCAAATGGGATCCATTCCAGTTGTTGGAAATGTTCCCAACCTAGTATCTTCT | 605 | | | | | | |
| QY | 655 | GTTCCCAACAGCAGCTGTGCCCTCCCTGCTAACCGGGCTCCCTCTGTTATACAACTCTG | 714 | | | | | | |
| DB | 606 | GTTCCCAACAGCAGCTGTGCCCTCCCTGCTAACCGGGCTCCCTCTGTTATACAACTCTG | 665 | | | | | | |
| QY | 715 | CCTGCATTTGCTCATCTCGAGCCACAATTCGCAAGAGTTCTTCTTTAGTAGATCTGGT | 774 | | | | | | |
| DB | 666 | CCTGCATTTGCTCATCTCGAGCCACAATTCGCAAGAGTTCTTCTTTAGTAGATCTGGT | 725 | | | | | | |
| QY | 775 | CCAGGCTCACAACCTAAACTAAATACAAAGGCACAGTCATTTGATGTGGCCAGTTC | 834 | | | | | | |

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|----|------|---|------|
| DB | 726 | CCAGGGTCACAACTAAACACTAAATTA CAAAAGGCACAGTCATTTGATGTGCCAGTGTCT | 785 |
| QY | 835 | CCACCAGTGGCAGAGTGGGCTGTTCTCAGTCATCAAGACTGAAATATACAGGCAATATTCT | 894 |
| DB | 786 | CCACCAGTGGCAGAGTGGGCTGTTCTCAGTCATCAAGACTGAAATATACAGGCAATATTCT | 845 |
| QY | 895 | AATAGTCATGACAAAACTATGAGTGGACACTTTAAC-----AGTCTCCCAAGCA | 942 |
| DB | 846 | AATAGTCATGACAAAACTATGAGTGGACACTTTAACAGAGTTCCTGTTTAGTCTCCCAAGCA | 905 |
| QY | 943 | AGAACTATTCTTATGAGTCAAGTTTACCAAGGCTCAGCTGGCTTCAATATGGAATCTT | 1002 |
| DB | 906 | AGAACTATTCTTATGAGTCAAGTTTACCAAGGCTCAGCTGGCTTCAATATGGAATCTT | 965 |
| QY | 1003 | TCTGCATTTGATCAAGATGGAATACTTACAGCAGAGGAAATTTATCTGGCAATGCACTTC | 1062 |
| DB | 966 | TCTGCATTTGATCAAGATGGAATACTTACAGCAGAGGAAATTTATCTGGCAATGCACTTC | 1025 |
| QY | 1063 | ATTGATGTAGTATGTTCTGGCCAAACCACTGCCACCTGTCCTGCTCCAGAAATCATTTCCA | 1122 |
| DB | 1026 | ATTGATGTAGTATGTTCTGGCCAAACCACTGCCACCTGTCCTGCTCCAGAAATCATTTCCA | 1085 |
| QY | 1123 | CTTCTTTTGAAGAGTTCGATCTGCGAGTGTATATCTGTCTATAGACTCAACATCTGTA | 1182 |
| DB | 1086 | CTTCTTTTGAAGAGTTCGATCTGCGAGTGTATATCTGTCTATAGACTCAACATCTGTA | 1145 |
| QY | 1183 | GATCAGAGGCTACCAAGAGAACCCAGTTTGTAGAAAGATGAAACAAACAATTTAGAAAAAGAA | 1242 |
| DB | 1146 | GATCAGAGGCTACCAAGAGAACCCAGTTTGTAGAAAGATGAAACAAACAATTTAGAAAAAGAA | 1205 |
| QY | 1243 | TTACTCTGTAACGTTTGAAGATGAAGACGGGAGAACTTTGAACTGGCAACTCTGGAACTG | 1302 |
| DB | 1206 | TTACTCTGTAACGTTTGAAGATGAAGACGGGAGAACTTTGAACTGGCAACTCTGGAACTG | 1265 |
| QY | 1303 | GAGAAACGAAAGGCAAGCTCTCTGGAACAGCAGCAGCAAGAGCAGGAGCCCTGGCCAG | 1362 |
| DB | 1266 | GAGAAACGAAAGGCAAGCTCTCTGGAACAGCAGCAGCAAGAGCAGGAGCCCTGGCCAG | 1325 |
| QY | 1363 | CTGGAGCGGCGGAGCAG | 1422 |
| DB | 1326 | CTGGAGCGGCGGAGCAG | 1385 |
| QY | 1423 | CAACTGGAACCTGGAGAGCAACTGGAAAGCAGCGGAGCTAGAAACGCGCAGAGAGAGAG | 1482 |
| DB | 1386 | CAACTGGAACCTGGAGAGCAACTGGAAAGCAGCGGAGCTAGAAACGCGCAGAGAGAGAG | 1445 |
| QY | 1483 | GAGAGAGGAGAAAGAAATTTGAGAGCGCAGAGAGCTGCAAAACCGGAACTTTGAAAGCAACGA | 1542 |
| DB | 1446 | GAGAGAGGAGAAAGAAATTTGAGAGCGCAGAGAGCTGCAAAACCGGAACTTTGAAAGCAACGA | 1505 |
| QY | 1543 | CAACTGAGTGGGAAACGGAATCGAAGCAAGAACTACTAAATCAAGAAACAAAGAAACAA | 1602 |
| DB | 1506 | CAACTGAGTGGGAAACGGAATCGAAGCAAGAACTACTAAATCAAGAAACAAAGAAACAA | 1565 |
| QY | 1603 | GAGGACATAGTTGTTACTGAAAGCAAAAGAAAGAACTTTGAAATTTGAAATTTAGAACTCTA | 1662 |
| DB | 1566 | GAGGACATAGTTGTTACTGAAAGCAAAAGAAAGAACTTTGAAATTTGAAATTTAGAACTCTA | 1625 |
| QY | 1663 | AATGATATAAAGCAATCAACTAGAGGAGAACTTCAAGATATCAGATCTCGATTGACCAACC | 1722 |
| DB | 1626 | AATGATATAAAGCAATCAACTAGAGGAGAACTTCAAGATATCAGATCTCGATTGACCAACC | 1685 |
| QY | 1723 | CAAGGCAAGAAATTTGAGAGCACAACAAATCTAGAGAGTTGAGAAATTTGCCGAATCAACC | 1782 |
| DB | 1686 | CAAGGCAAGAAATTTGAGAGCACAACAAATCTAGAGAGTTGAGAAATTTGCCGAATCAACC | 1745 |
| QY | 1783 | CATCTACAGCAACAAATTTACAGGAATCTCAGCAATGCTTGGAAAGCTTTATTTCCAGAAAA | 1842 |
| DB | 1746 | CATCTACAGCAACAAATTTACAGGAATCTCAGCAATGCTTGGAAAGCTTTATTTCCAGAAAA | 1805 |
| QY | 1843 | CAGATACCTCAATGACCAATTTAAACAAAGTTTCAGCAGAACAGTTTGGCAGAGATTTCATT | 1902 |
| DB | 1806 | CAGATACCTCAATGACCAATTTAAACAAAGTTTCAGCAGAACAGTTTGGCAGAGATTTCATT | 1865 |

1903 GTTACACTTAAAGAGCCCTTAGAAGCAAAAGAACTAGCTCGCAGACCACTACGAGACCAA 1962
1866 GTTACACTTAAAGAGCCCTTAGAAGCAAAAGAACTAGCTCGCAGACCACTACGAGACCAA 1925
1963 CTGGATGAAGTGGAGAAAGAACTAGATCAAACTACAGGAGTTGATATTTTCATAT 2022
1926 CTGGATGAAGTGGAGAAAGAACTAGATCAAACTACAGGAGTTGATATTTTCATAT 1985
2023 CAGCTGAAGCACTAAGAGAAATACAAATAAGCAAACTCCAGAGCAAAAGTCCATG 2082
1986 CAGCTGAAGCACTAAGAGAAATACAAATAAGCAAACTCCAGAGCAAAAGTCCATG 2045
2083 GAGGCTGAACGACTGAAACAGAAAGAAACGAAAGATCATAGAAATAGAAAACAA 2142
2046 GAGGCTGAACGACTGAAACAGAAAGAAACGAAAGATCATAGAAATAGAAAACAA 2105
2143 AAAGAAGAGCCCAAGACGAGCTCAGGAAAGGGACAAAGCTGGCTGGAGCATGTGCAG 2202
2106 AAAGAAGAGCCCAAGACGAGCTCAGGAAAGGGACAAAGCTGGCTGGAGCATGTGCAG 2165
2203 CAGGAGGACGAGCATCAGAGACCAAGAAAACCTCCAGAAAGAGGAAAACTGAAAAGGGAG 2262
2166 CAGGAGGACGAGCATCAGAGACCAAGAAAACCTCCAGAAAGAGGAAAACTGAAAAGGGAG 2225
2263 GAGAGTGTCAAAAGAGGATGGCGAGGAAAAAGGCAAAACAGGAAGCAACAAGCTG 2322
2226 GAGAGTGTCAAAAGAGGATGGCGAGGAAAAAGGCAAAACAGGAAGCAACAAGCTG 2285
2323 GGTGGCTTTTCATCAACACCAAGAACGAGCTTAGCCAGCTGTCCAGGCACCTGGTCC 2382
2286 GGTGGCTTTTCATCAACACCAAGAACGAGCTTAGCCAGCTGTCCAGGCACCTGGTCC 2345
2383 ACTGCAGAAAAAGGTCCACTTACCATTTCTGCACAGGAAAAATGTAAAGTGGTGATTAC 2442
2346 ACTGCAGAAAAAGGTCCACTTACCATTTCTGCACAGGAAAAATGTAAAGTGGTGATTAC 2405
2443 CGGGCACTGTACCCCTTTGAATCCAGAGCCATGATGAATCACTATCCAGCCAGAGAG 2502
2406 CGGGCACTGTACCCCTTTGAATCCAGAGCCATGATGAATCACTATCCAGCCAGAGAG 2465
2503 ATAGTCATGGTGGATGAAAGCCAACTGGAGAACCCGGCTGGCTTGGAGGAGAAATTA 2562
2466 ATAGTCATGGTGGATGAAAGCCAACTGGAGAACCCGGCTGGCTTGGAGGAGAAATTA 2525
2563 GGAAAGACAGGGTGGTTCCCTGCAAACTATGCAGAGAAATCCAGAAATGAGGTTCCT 2622
2526 GGAAAGACAGGGTGGTTCCCTGCAAACTATGCAGAGAAATCCAGAAATGAGGTTCCT 2585
2623 GCTCCAGTGAACCAAGTGAATCAACATCTGCCCCCTGCCCCCAAACTGGCCTTGGCT 2682
2586 GCTCCAGTGAACCAAGTGAATCAACATCTGCCCCCTGCCCCCAAACTGGCCTTGGCT 2645
2683 GAGACCCCGCCCTTTGGCAGTAACTCTTCCAGAGCCCTCCAGACCCCTTAATACCTG 2742
2646 GAGACCCCGCCCTTTGGCAGTAACTCTTCCAGAGCCCTCCAGACCCCTTAATACCTG 2705
2743 GCCGACTTCAGTTCACGCTGGCCCAACAGCAAGATGAGAAAACGAGAACTGATCTGG 2802
2706 GCCGACTTCAGTTCACGCTGGCCCAACAGCAAGATGAGAAAACGAGAACTGATCTGG 2765
2803 GATGATGGGACCCAGCCCTCTCTCAGCGTTCCAGTGGCGGCCAGTAAAGGACAGAG 2862
2766 GATGATGGGACCCAGCCCTCTCTCAGCGTTCCAGTGGCGGCCAGTAAAGGACAGAG 2825
2863 TCCGCTTTACTCCAGCAGCCACTGGCTCTCCCGTCTCTCTGCTAGGCCAGGCT 2922
2826 TCCGCTTTACTCCAGCAGCCACTGGCTCTCCCGTCTCTCTGCTAGGCCAGGCT 2885
2923 GA 2924
2886 GA 2887

RESULT 7
US-10-264-049-887
; Sequence 887, Application US/10264049
; Publication No. US20040005579A1
; GENERAL INFORMATION:
; APPLICANT: Birse et al.
; TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
; FILE REFERENCE: PA133PI
; CURRENT APPLICATION NUMBER: US/10/264,049
; PCT FILING DATE: 2002-10-04
; PRIOR APPLICATION NUMBER: PCT/US01/18569
; PRIOR FILING DATE: 2001-06-07
; PRIOR APPLICATION NUMBER: US 60/209,467
; PRIOR FILING DATE: 2000-06-07
; NUMBER OF SEQ ID NOS: 4360
; SOFTWARE: PatentIn Ver. 3.1
; SEQ ID NO 887
; LENGTH: 2067
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (5)-(5)
; OTHER INFORMATION: n equals a,t,g, or c
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (2058)-(2058)
; OTHER INFORMATION: n equals a,t,g, or c
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (2063)-(2063)
; OTHER INFORMATION: n equals a,t,g, or c
US-10-264-049-887

Query Match 36.2%; Score 1882.4; DB 6; Length 2067;
Best Local Similarity 98.3%; Pred No. 0;
Matches 1953; Conservative 1; Mismatches 27; Indels 5; Gaps 5;

QY 3199 TCGGAGAGAAATTTGCCAGGTTATTCCTCATACACCGCACCGGGCCCCGAGCAGCTC 3258
Db 53 TCTCAAAATAAATTTCCCCAGGTTATTGCCTCATACACCGCACCGGGCCCCGAGCAGCTC 112
QY 3259 ACTCTCGCCCTGGTCAGCTGATTTTGTATCCGAAAAAGAACCCAGGTGGATGGGAA 3318
Db 113 ACTCTCGCCCTGGTCAGCTGATTTTGTATCCGAAAAAGAACCCAGGTGGATGGGAA 172
QY 3319 GGAGAGCTGCAAGCACGTGGGAAAAAGCGCCAGATAGGCTGTTCCAGCTAATTTATGTA 3378
Db 173 GGAGAGCTGCAAGCACGTGGGAAAAAGCGCCAGATAGGCTGTTCCAGCTAATTTATGTA 232
QY 3379 AAGCTTTAAGCCCTGGGACGAGCAAAATCACTCCAAACAGACCCACCTTAAGTCAACAGCA 3438
Db 233 AAGCTTTAAGCCCTGGGACGAGCAAAATCACTCCAAACAGACCCACCTTAAGTCAACAGCA 292
QY 3439 TTAGCGGCAAGTGTGCGCAGGTGATTTGGGATGTACGACTACACCGCCAGAAATGCGATGAG 3498
Db 293 TTAGCGGCAAGTGTGCGCAGGTGATTTGGGATGTACGACTACACCGCCAGAAATGCGATGAG 352
QY 3499 CTGGCTTCAACAAAGGGCCAGATCATCAACGTCTTCAACAGGAGGAGCCCTGACTGGTGG 3558
Db 353 CTGGCTTCAACAAAGGGCCAGATCATCAACGTCTTCAACAGGAGGAGCCCTGACTGGTGG 412
QY 3559 AAAGAGAGAGTCAATGGACAAAGTGGGCTTTTCCCATCAATATGTGAAGCTGACCAACA 3618
Db 413 AAAGAGAGAGTCAATGGACAAAGTGGGCTTTTCCCATCAATATGTGAAGCTGACCAACA 472
QY 3619 GACATGGACCCCAAGCCCAATGAATCATATGTTGTCATATCCCTCCAGCTTCAAGAG 3678
Db 473 GACATGGACCCCAAGCCCAATGAATCATATGTTGTCATATCCCTCCAGCTTCAAGAG 532
QY 3679 TCTTCAAGAGAGACCCCACTATCCCATATCACTGCCAGAGGGGATGATGGAGATGACGCTT 3738

[illegible]

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RESULT 8
US-10-450-763-20566
; Sequence 20566, Application US/10450763
; Publication No. US20050196754A1
; GENERAL INFORMATION:
; APPLICANT: Hveeq. Inc
; TITLE OF INVENTION: NOVEL NUCLEIC ACIDS AND POLYPEPTIDES
; FILE REFERENCE: 790CIP3/US
; CURRENT APPLICATION NUMBER: US/10/450,763
; CURRENT FILING DATE: 2003-06-11
; PRIOR APPLICATION NUMBER: PCT/US01/08631
; PRIOR FILING DATE: 2001-03-30
; PRIOR APPLICATION NUMBER: 09/540,217
; PRIOR FILING DATE: 2000-03-31
; PRIOR APPLICATION NUMBER: 09/649,167
; PRIOR FILING DATE: 2000-08-23
; NUMBER OF SEQ ID NOS: 60736
; SOFTWARE: Custom
; SEQ ID NO 20566
; LENGTH: 2874
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: SIMILAR
; LOCATION: (1)..(1545)
; OTHER INFORMATION: 100% homologous to Homo sapiens Human SH3D1A
; OTHER INFORMATION: protein,accession number Y32156,Smith-Waterman
US-10-450-763-20566
Score=2669.

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| Qy | 1513 | GCTGCAAAACGGGAATCTTGAAAGGCAACGACAACTTGAGTGGGAAACGGAATCGAAGCAA | 1572 |
| Db | 1 | GCTGCAAAACGGGAATCTTGAAAGGCAACGACAACTTGAGTGGGAAACGGAATCGAAGCAA | 60 |
| Qy | 1573 | GAACTACTTAATCAAAAGAAAACAAAGAAACAAAGAGGACATAGTTGTACTGAAAGCAAAGAAA | 1632 |
| Db | 61 | GAACTACTTAATCAAGAAACAAAGAAACAAAGAGGACATAGTTGTACTGAAAGCAAAGAAA | 120 |
| Qy | 1633 | AAGACTTTGGAAATTTGAAATTAGAGCTCTAAATGATATAAAGCATCAACTTAGAGGGAAA | 1692 |

Db 121 AAGACTTTGGAAATTTGAATTTAGAGCTCTAAATGATATAAAAAAGCATCAACTAGAGGGAAA 180
Qy 1693 CTTTCAAGATATCAGATGTGCGATTGACCAACCCAAAGGCAAGAAATTTGAGAGCACAACCAA 1752
Db 181 CTTTCAAGATATCAGATGTGCGATTGACCAACCCAAAGGCAAGAAATTTGAGAGCACAACCAA 240
Qy 1753 TCTAGAGAGTTGAGAAATTTGCCGAATCACCATCTTACAGCAACAATTTACAGGAATCTCAG 1812
Db 241 TCTAGAGAGTTGAGAAATTTGCCGAATCACCATCTTACAGCAACAATTTACAGGAATCTCAG 300
Qy 1813 CAAATGCTTTGGAAGACTTATTTCCAGAAAAACAGATCTCAATGACCAATTTAAAAACAAGTT 1872
Db 301 CAAATGCTTTGGAAGACTTATTTCCAGAAAAACAGATCTCAATGACCAATTTAAAAACAAGTT 360
Qy 1873 CAGCAGAAACAGTTTGCACAGAGATTCACCTTTGTGTACACTTTAAAGAGCCTTTAGAAGCAAAA 1932
Db 361 CAGCAGAAACAGTTTGCACAGAGATTCACCTTTGTGTACACTTTAAAGAGCCTTTAGAAGCAAAA 420
Qy 1933 GAACTAGCTCGGAGCAGCACTACAGACCAACTGGATGAAGTGGAGAAAGAACTAGATCA 1992
Db 421 GAACTAGCTCGGAGCAGCAGCACTACAGACCAACTGGATGAAGTGGAGAAAGAACTAGATCA 480
Qy 1993 AAATCTACAGAGATTTGATATTTTCAATAATCAGCTGAAGGAACTAAGAGAAATACACAAT 2052
Db 481 AAATCTACAGAGATTTGATATTTTCAATAATCAGCTGAAGGAACTAAGAGAAATACACAAT 540
Qy 2053 AAGCAACAACTCCAGAAAGCAAAAGTCCATGGAGGCTGAAACGACTGAAACAGAAAGAACAA 2112
Db 541 AAGCAACAACTCCAGAAAGCAAAAGTCCATGGAGGCTGAAACGACTGAAACAGAAAGAACAA 600
Qy 2113 GAAAGCAAGATCATGAAATTTAGAAAAAACAAGAAAGAACCCAAAGACAGCTCAGGAA 2172
Db 601 GAAAGCAAGATCATGAAATTTAGAAAAAACAAGAAAGAACCCAAAGACAGCTCAGGAA 660
Qy 2173 AGGGCAACAGCTGGCTGAGCATGTGCAGAGGAGGAGAGCATCAGAGACCAAGAAA 2232
Db 661 AGGGCAACAGCTGGCTGAGCATGTGCAGAGGAGGAGAGCATCAGAGACCAAGAAA 720
Qy 2233 CTTCCAGAAAGAGGAAAACTGAAAAGGGAGGAGGTGTCAAAAAGAAAGGATCGCAGGAA 2292
Db 721 CTTCCAGAAAGAGGAAAACTGAAAAGGGAGGAGGTGTCAAAAAGAAAGGATCGCAGGAA 780
Qy 2293 AAAGGCAACAGAGGACCAAGACAGCTGGGTGCGCTTTTCCATCAACACCAAGAACCA 2352
Db 781 AAAGGCAACAGAGGACCAAGACAGCTGGGTGCGCTTTTCCATCAACACCAAGAACCA 840
Qy 2353 GCTAAGCCAGCTGTCCAGGACCCCTGGTCCACTGCAGAAAAAGGTCCACTTACCATTCT 2412
Db 841 GCTAAGCCAGCTGTCCAGGACCCCTGGTCCACTGCAGAAAAAGGTCCACTTACCATTCT 900
Qy 2413 GCACAGGAAAAATGTAAAAGTGTGTATTACCGGGCACTGTACCCCTTTGAAATCCAGAAAC 2472
Db 901 GCACAGGAAAAATGTAAAAGTGTGTATTACCGGGCACTGTACCCCTTTGAAATCCAGAAAC 960
Qy 2473 CATGATGAAATCACTATCAGCCAGAGACATAGTTCATGGTGGATGAAGCCAAACTGGA 2532
Db 961 CATGATGAAATCACTATCAGCCAGAGACATAGTTCATGGTGGATGAAGCCAAACTGGA 1020
Qy 2533 GAAACCGGCTGGCTTCGAGAGAAATTTAAAGGAAAGACAGGGTGTTCCTCGCAAACTAT 2592
Db 1021 GAAACCGGCTGGCTTCGAGAGAAATTTAAAGGAAAGACAGGGTGTTCCTCGCAAACTAT 1080
Qy 2593 GCAGAGAAAAATCCAGAAAAATGAGTTCCCGCTCCAGTGAACCAGTGTGATTTCAACA 2652
Db 1081 GCAGAGAAAAATCCAGAAAAATGAGTTCCCGCTCCAGTGAACCAGTGTGATTTCAACA 1140
Qy 2653 TCTGCCCTTCCCCGAACTGGCTTGGCTGAGACCCCGCCCTTTTGGCAGTTACCTCT 2712
Db 1141 TCTGCCCTTCCCCGAACTGGCTTGGCTGAGACCCCGCCCTTTTGGCAGTTACCTCT 1200
Qy 2713 TCAGAGCCCTCCACGACCCCTTAATAACTGGGCGGACTTCAGCTCCACGTGGCCACACG 2772
Db 1201 TCAGAGCCCTCCACGACCCCTTAATAACTGGGCGGACTTCAGCTCCACGTGGCCACACG 1260

Qy 2773 ACGAATGAGAAACCGAGAAACCGATAACTGGGATGATGGGAGCCCGCCCTCTCTCACC 2832
Db 1261 ACGAATGAGAAACCGAGAAACCGATAACTGGGATGATGGGAGCCCGCCCTCTCTCACC 1320
Qy 2833 GTTCCAAGTGGCGGCGAGTTAAGGCAGAGTCCGCTTTTACTTCCAGCCACCGCCACTGGC 2892
Db 1321 GTTCCAAGTGGCGGCGAGTTAAGGCAGAGTCCGCTTTTACTTCCAGCCACCGCCACTGGC 1380
Qy 2893 TCCTCCCGCTCTCTCTGTGTAGGCTAGGCTGAAAGAGTGGAGGGCTACAAGCTCAAGC 2952
Db 1381 TCCTCCCGCTCTCTCTGTGTAGGCTAGGCTGAAAGAGTGGAGGGCTACAAGCTCAAGC 1440
Qy 2953 CTATATCTTGGAGAGCCAAAAAGACAAACACATTAATTTTAAACAAAAATGATCTCATC 3012
Db 1441 CTATATCTTGGAGAGCCAAAAAGACAAACACATTAATTTTAAACAAAAATGATCTCATC 1500
Qy 3013 ACCGTCTCGGAACAGCAAGACATGTGTGGTGGTGGAGAGTTCAAGGTCAAGAGGTTGG 3072
Db 1501 ACCGTCTCGGAACAGCAAGACATGTGTGGTGGTGGAGAGTTCAAGGTCAAGAGGTTGG 1545
Qy 3073 TTCCCAAGTCTTACGTGAAACTCATTTTCAAGGCCCATTAAGGAAGTCTACAAGCATGAT 3132
Db 1546 -----GAAATTTATGGCATGTACACTTACGAGAGTTCTTGAGCAAGGAG 1588
Qy 3133 TCTGTTCTTTCAGAGAGTCTCTGTAGTCTAAGCGAGTAGCTCTCAGCAGCACAAGCGG 3192
Db 1589 ATTTAACCTT-----TCAAGTCTTAAAGATTTCAGAGGGCTCTGGAA 1605
Qy 3193 GTCTGTTTCGGGAGAGAAATTTGCCAGGTATTGGCTCATACACCGCCACCGGCCCGGAG 3252
Db 1606 GGGGATGTATTTTGGTTTACCAGAAAGATGTGTGACTGTGGACAGGAAACAGTGGCGGAC 1665
Qy 3253 CAGCTCACTCTCGCCCTCGTGTGAGTGTGATTCGAAAGAAAGAAACCCAGGTGGATGG 3312
Db 1666 AAGCCGCGAGTCTTCCCT--TCTAACTATGTGAGGCTTAAAGATTTCAGAGGGCTCTGGAA 1723
Qy 3313 TGGGAAGAGAGAGTCTCAAGCAGCTGGGAAAGCCAGATAGGTGGTTCACAGTAAAT 3372
Db 1724 CTGCTGGGAAACACAGGGAGTTTAGGAAAGAAACCTGAAATTTGCCAGGTATTGGCTCAT 1783
Qy 3373 TATGTAAAGCTTCTAAGCCCTGGGACGAGCAAAATCACTCCAAACAGAGCCACCTAAGTCA 3432
Db 1784 ----ACACCGCCACCGGCCCGGAGCAGCTCACTCTCGCCCTTGGTCAAGTATTGATC 1839
Qy 3433 ACAGCATTTAGCGGCAAGTGTGCAGGTGATGGGATGTACGACTACACCGCGCAGAAATGAC 3492
Db 1840 CGAAAAAGAAACCCAGTGTGCCAGGTGATGGGATGTACGACTACACCGCGCAGAAATGAC 1899
Qy 3493 GATGAGCTGGCTTCAACAGGGCCAGATCATCAAGCTCTCAACAGGAGGACCCCTGAC 3552
Db 1900 GATGAGCTGGCTTCAACAGGGCCAGATCATCAACGCTCTCAACAGGAGGACCCCTGAC 1959
Qy 3553 TGGTGGAAAGGAGAAAGTCAATGGCAAGTGGGGCTCTTCCCATCCAAATTTATGTGAAGCTG 3612
Db 1960 TGGTGGAAAGGAGAAAGTCAATGGCAAGTGGGGCTCTTCCCATCCAAATTTATGTGAAGCTG 2019
Qy 3613 ACCCAGACATGGAGCCCAAGCCAGCAAT 3640
Db 2020 ACCCAGACATGGAGCCCAAGCCAGCAAT 2047

RESULT 9

US-10-398-885A-15
; Sequence 15, Application US/10398885A
; Publication No. US20040053282A1
; GENERAL INFORMATION:
; APPLICANT: Sugita, Yuji
; APPLICANT: Hashida, Ryoichi
; APPLICANT: Ogawa, Kaoru
; APPLICANT: Nagasu, Takashi
; APPLICANT: Obayashi, Masaya
; APPLICANT: Saito, Hirohisa


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QY 1918 GCCTTAGAGCAAAAGAACTAGCTGGCAGACCTACGAGACCAACTGATGAAGTGGAG 1977
Db 1789 AAATCATTTAGAAAAGCAAAATTAATGCCAAAGACTTAAAGAACAGTTAGATGCTCTTGA 1848
QY 1978 AAAGAACTAGATCAAACTACAGGAGATTGATATTTCAATAATCAGCTGAAGGACTA 2037
Db 1849 AAAGAAACTGCATCTAAGCTGTAGAAATGGATTCCTTTAAACAATCAACTAAAGGAATG 1908
QY 2038 AGAGAAATACAAATTAAGCAAACTCCAGAAAGCAAAAGTCCATGAGGCTGAAAGCACTG 2097
Db 1909 AGAGAAACCTACAAACACAGCAGTTAGCCCTTGAACACTTTATAAGATCAAACTGTAC 1968
QY 2098 AAACAGAAAGAACAGAAAGCAAGATCATAGAAAT-----AGAAAAACAAAGAA 2148
Db 1969 AAGTTGAAGGAAATTTGAAAGGAAAGATTTAGAACTAATGCAGAAAAAGAAACTTAGAAT 2028
QY 2149 GAAGCCAAAGCAGAGCTCAGGAAAGGCAAGCAGTGTCTGGAGCATGTGCAGCAGGAG 2208
Db 2029 GAGGCTGCAAGGAAAGCAAGCAAGCAAGGAAAGAAACTTATGAAAGAAAAATCTTAGAAAG 2088
QY 2209 GACGAGCATCAGAGACCAAGAAAACTCCACGAAGAGGAAAAAAGTGAAGGGAGGAGAGT 2268
Db 2089 GAGGAGAGAAAAACAAAGCGACTCCAGGAAAGAAAAACACAGAAAAAATTCAGAA 2148
QY 2269 GTCAAAAGAGAGATGGCGAGGAAAAAGGCAACAGGAAG-CACAAAGCAAGCTGGTGTG 2327
Db 2149 GAGGAACGGAAGCTGAGGAGAAAAACAACGTAAGGATAAGGATACTTTTGAAGCTGAGGAG 2208
QY 2328 GCTTTTCCATCAACACCAAGAACAGCTAAGCCAGCTGTCCAGGCACCTGTCTCACTGC 2387
Db 2209 AAAAAACGTGAGACAGCTAGTGTGTTTGGTGAATATATAGAGCATATATACCCCTTTGA 2268
QY 2388 AGAAAAAGGTCACCTTACCATTCTGCACAGAAAAATGTAAAGTGGTGTATTTACCGGC 2447
Db 2269 AGGAACCATGATGAGATGAGTTTAAATCTGGAGATATAATTCAGGTGATGAAAAAAC 2328
QY 2448 ACTGTACCCCTTTGAATCCAGAGCCATGATGAATCATATCCAGCCAGGAGACATAGT 2507
Db 2329 GTAGGAGAACTGGTTGGCTTTATGTTAGTGTTCGAAGGAAATTTTGGCTGGTTTCATGC 2388
QY 2508 CATGGTGGATG-----AGCCAACTGGAGAACCCGGCTGGCTTGGAGGAA 2556
Db 2389 AATTATGTAGAAAAAATGCCATCAAGTGAATAATGAAAAAGCTGTATCTCAAAGAGGCC 2448
QY 2557 TTAAGAGGAAAGACAGGGTGGTTCCCTGCAAACT---ATGCAGAGAAAAATCCAGAAAA 2613
Db 2449 TTACTTCCTCTACAGTTCTTTATCTGTACTCACTTCCCTCTGAAACCACTTTCTTCA 2508
QY 2614 GAGGTTCCGCTCCAGTGAACCAAGTGAATCAACATCTGCCCCCTGCCCCCAAACTG 2673
Db 2509 AATCAACCAAGCATCAGTCACTGATTTATCAAAATGTATCTTTTCAAACCTAACTGTAA 2568
QY 2674 GCCTTGGGTGAGACCCCGCCCTTTGGCAGTAACTCTTCAGAGCCCTCCAGACCCCT 2733
Db 2569 ACATCATGCGAGAAAAAATCAGCCTTCACTCGAACTGTGTCCCTGGATCTGTATCACCT 2628
QY 2734 AATAACTGG-----GCCACTTTCAGCTCCACCTGCGCCACCCAGCAGCAATGAGAAA 2784
Db 2629 ATTCATGGACAGGCAAGTGGTAGAAAACCTTAAAGACACAGGCCCTTTGTTCTTGACT 2688
QY 2785 CCAGAAACCGATAACTGGGATGATGGGAGCCAGCCCTCTCTCACCGTTCCAAAGTGGC 2844
Db 2689 GCAAGAAAAAGATAACCACTTGAATCTCAAAAACATGACATATTACTGTCTTTGGAGCAG 2748
QY 2845 GGCCAGTTAAGCAGAGGTCGCGCTTTACTCCAGCCACGGCCACTGGCTCCTCCCGTCT 2904
Db 2749 CAAGAAAAATTTGGTGGTGGGAGGTGCATGGAGGAAGAGGATGGTTTCCCAAAATCTTAT 2808
QY 2905 CTGTGCTAGGCGAGGGTGAAAAAGGTGGAGGGGCTACAAGCTCAAGCCCTTATCTCTGG 2964
Db 2809 GTCAAGATCATCTCTGGAGTGAAGTAAAAACGGGAGAACCAAGAACTTTGTATGAGCT 2868
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QY 2965 AGAGCCAAAAAGACCAACCACTTAAATTTTAAACAAAATGATGTATCACCGTCTCTGGAA 3024
Db 2869 GTAAATTAAGAAACCTTACCTCGCAGCCTATTTCAGTTGGAGAGAAATATATTGCACTTTAT 2928
QY 3025 CAGCAAGACATGTGTGTGGTGGTGGAGAGTTCAAGGTGAGAAAGGGTTGGTTCCCCAAGTCT 3084
Db 2929 C-CATATTCAAGTGTGGAACTCTGGAGATTTCAGCTTTTACAGAGGTGAAGAAATATTGGT 2987
QY 3085 TACGTGAAACTCATTTTCAGGGCCCATAAAGGAAGTCTACAAGCATGGATTCTGGTTCTTCA 3144
Db 2988 GACCCAGAAAAGATGGAGAGTGTGGACAGGAAGTATTGGAGATAGAAGTGGAAATTTTTCC 3047
QY 3145 GAGAGTCTGTCT-----AGTCTAAAGCCAGTAGCCTCTCCACGACCCAGCCGCTCG 3196
Db 3048 ATCAAACTATGTCAAAACCAAGAGGATCAAGAGAGTTTGGGAGTGTAGCAAGTCTGGAGC 3107
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Db 3108 ATCAATTAATAAAACCTGAGATTGCTCAGGTAACTTCAGCATATGTTGCTTCTGGTTCTGA 3167
QY 3252 GCAGTCACTCTCGCCCTGTGTCAGTCAATTTTGTATCCGAAAAAGAACCCAGGTGGATG 3311
Db 3168 ACAACTTAGCCTTGCACCGACAGTTAATTAATTTCTAAAGAAAAATACAAGTGGGTG 3227
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Db 3228 GTGGCAAGGAGAGTTACAGGCCAGAGGAAAGGCGACAGAAAGGATGGTTTCTGCGCAG 3287
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Db 3288 TCATGTTAACTTTTGGGTCCAAAGTAGTGAAGAGCCA-----C 3326
QY 3432 AACACATTAGCGGAGTGTGCCAGTGTGCGAGTGAATTTGGAGTATGACACTACACCGCCAGAA 3491
Db 3327 ACCTGCCCTTCATCTGTATGTGTCAGGTGATGCTATGATGACTATGACTATGAGCAAAATGA 3386
QY 3492 CGATGAGCTGGCTTCAACAAAGGCCAGATCATCAAGCTCTCAACAAAGGAGGACCTGA 3551
Db 3387 AGATGAGCTCAGTTTCTCAAAGGGCAACTCATTAATGTTATGAAAGAGATGATCTCTGA 3446
QY 3552 CTGTTGGAAGAGAGAAAGTCAATGGACAAGTGGGGCTTCTTCCCATCCAATTTATGTAAGCT 3611
Db 3447 TTGTTGGCAAGGAGAGATCAACGGGTGACTGGTCTCTTCTTCCCTCAAACCTAGTTAAGAT 3506
QY 3612 GACACAGACATGGACCCCAAGCCAGCAATG 3641
Db 3507 GACGACAGACTCAGATCCAAGTCAACAGTG 3536
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RESULT 10

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US-09-884-441-72
; Sequence 72, Application US/09884441
; Patent No. US20020119158A1
; GENERAL INFORMATION:
; APPLICANT: Algate, Paul A.
; APPLICANT: Carcer, Darrick
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
; TITLE OF INVENTION: DIAGNOSIS OF OVARIAN CANCER
; FILE REFERENCE: 210121.462C7
; CURRENT APPLICATION NUMBER: US/09/884,441
; CURRENT FILING DATE: 2001-06-18
; NUMBER OF SEQ ID NOS: 489
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 72
; LENGTH: 2017
; TYPE: DNA
; ORGANISM: Homo sapien
US-09-884-441-72
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Query Match 9.8%; Score 507.8; DB 3; Length 1017;
Best Local Similarity 55.8%; Pred. No. 4.9e-107;
Matches 1138; Conservative 0; Mismatches 802; Indels 99; Gaps 5;

| | | | |
|----|------|---|------|
| QY | 175 | GGCGTCGATTAGCAAGGTAAGTAACAGAACCAATCGCTCAGTTTCCACACCTTTGGT | 234 |
| Db | 9 | GGCTGAGAGCTGCAGAAAGAGTCAGGATCATGATGGCTCAGTTTCCACAGGATGAAT | 68 |
| QY | 235 | GGCAGCCTGGATATCTGGGCCATAACTGTAGAGGAAGAGCGAAGCATGATCAGCAGTTTC | 294 |
| Db | 69 | GGAGGCCAATATGTGGGCTATTACATCTGAAGACGACTTAAGCATGATAAACAAGTTT | 128 |
| QY | 295 | CATAGTTTAAAGCCAAATATCTGGATTATTAATCTGTGGTATCAAGCTAGTAAGAACTTTT | 354 |
| Db | 129 | GATAACCTCAAACTTTCAGAGGTTACATAACAGGTGATCAAGCCCGTACTTTTTCCTA | 188 |
| QY | 355 | CAATCTGGGTACCTCAACTGTTTGTAGCAGATATGGCACTAGCTGACATGAATAAT | 414 |
| Db | 189 | CAGTCAAGGTCGCGGCCCGGTTTGTAGCTGAATATGGCCCTTATCAGATCTGAACAAG | 248 |
| QY | 415 | GATGGAAGATGATCAAGTGGAGTTTTCATAGCTATGAACCTTATCAAACTGAAGCTA | 474 |
| Db | 249 | GATGGGAAGATGGACAGCAAGATTTCTCTATAGCTATGAACCTCATCAAGTTAAAGTTG | 308 |
| QY | 475 | CAAGGATATCAGCTACCTCTGCACTTCCCTCTGTCTATGAACAGCAACG---AGTTGCT | 531 |
| Db | 309 | CAGGGCCAAAGCTGCTGTAGTCTCTCCCTCTATCATGAACCAACCCCTATGTTCTCT | 368 |
| QY | 532 | ATTTCTAGCCACCAGCATTTGGTATGGAGGTATCGCAGCATGCCACCGCTTACAGCT | 591 |
| Db | 369 | CCACTAATCTCTGCTCGTTTGGGATGGGAAGCATGCCCAATCTGTCCATTCATCAGCCA | 428 |
| QY | 592 | GTTCCTCCAGTGCATGGATC-----CATTT | 618 |
| Db | 429 | TTGCCCTCCAGTTGCATATAGCAACACCTTCTCTGCTACTTCAGGACCATGATTT | 488 |
| QY | 619 | CCAGTTGTTGGAATGTCTCAACCCCTAGTATCTTCTGTTCCCAAGAGCTGTGCCCCC | 678 |
| Db | 489 | CCTCCCTAATGATGCTGCTGCCCTAGTGCCTTCTGTTAGTACATCTCAITACCAAT | 548 |
| QY | 679 | CTGGCTAAAGGGGCTCCCTGTTATACAACTCTGCCTGCATTTGCTCATCTCGAGCC | 738 |
| Db | 549 | GGAACTGCCAGTCTCATTCAGCCTTTATCCATTCCTTATTTCTTCAACATTTGCCCTCAT | 608 |
| QY | 739 | ACATGTCGAAGAGTTCTTCTTTAGTAGATCTGGTCCAGGGTCACAACTTAAACACTAAA | 798 |
| Db | 609 | GCATCATCTTACGCTGATGATGGAGGATTTGGTGTCTAGTATCCAGAGGGCCAG | 668 |
| QY | 799 | TTACAAAGGCACAGTCATTTGATGTGGCCAGTGTCCACAG-----841 | 841 |
| Db | 669 | TCTCTGATTGATTAGGATCTAGTAGCTCAACTTCCTCAACTGCTTCCCTCTCAGGGAAC | 728 |
| QY | 842 | -----TGGCAGAGTGGCTGTTCTCAGTCATCAGACTGAAATACAGG | 885 |
| Db | 729 | TCACCTAAGACAGGACCTCAGAGTGGGCGAGTTCTCAGCCTTCAAGATTAAGATTCGG | 788 |
| QY | 886 | CAATATTCAATAGTATGACAAACTATGAGTGGACACTTAAACAGTCCCCCAAGCAAGA | 945 |
| Db | 789 | CAAAAATTTATAGTCTAGCAAAAGGATGAGCGGATACCTCTCAGGTTTCAAGCTAGA | 848 |
| QY | 946 | ACTATTCTTATGCACTCAAGTTTACACAGGCTCAGCTGGCTTCAATATGGAATCTTTCT | 1005 |
| Db | 849 | AATGCCCTTCTTCACTCAAACTCTCTCAAACTCAGCTAGCTACTATTTGGACTCTGGCT | 908 |
| QY | 1006 | GACATTCATCAGATGGAACACTTACAGCAGAGAAATTTATCTCGCAATGCACCTCATTT | 1065 |
| Db | 909 | GACATCGATGGTGCAGACAGTGTAAAGCTGAAGAAATTTATTTCTGGCGATGCACCTCACT | 968 |
| QY | 1066 | GATGTAGCTATGTCTGGCCACCACTGCACCTGCTCTGCTCCAGAAATATATTTCCACCT | 1125 |
| Db | 969 | GACATGGCCAAAGCTGGACACCACTACCACTGAGCTGTGCTCCGAGCTGTGCTCCCA | 1028 |
| QY | 1126 | TCTTTTAGAAAGTTCGATCTGGCAGTGGTATATCTGTCTATAGCTCAACATCTGTAGAT | 1185 |
| Db | 1029 | TCCTTCAGAGGGGAAAGCAAGTTGAT-----TCTGTTAATGGAACCTGCTCTCATAT | 1082 |
| QY | 1186 | CAGAGGCTACAGAGGACCACTTTTAGAAGATGAACAACTATAGAAAGAAATTA | 1245 |

RESULT 11
US-09-907-969-72
; Sequence 72, Application US/09907969
; Publication No. US20030091580A1
; GENERAL INFORMATION:
; APPLICANT: Mitcham, Jennifer L.
; APPLICANT: King, Gordon E.
; APPLICANT: Algate, Paul A.
; APPLICANT: Fling, Steven P.

```
/ APPLICANT: Retter, Marc W.
/ APPLICANT: Fanger, Gary Richard
/ APPLICANT: Reed, Steven G.
/ APPLICANT: Vedvick, Thomas S.
/ APPLICANT: Carter, Darrick
/ APPLICANT: Hill, Paul
/ APPLICANT: Albone, Earl
/ TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
/ TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
/ FILE REFERENCE: 210121.462C8
/ CURRENT APPLICATION NUMBER: US/09/907,969
/ CURRENT FILING DATE: 2001-07-17
/ NUMBER OF SEQ ID NOS: 596
/ SOFTWARE: FastSeq for Windows Version 4.0
/ SEQ ID NO 72
/ LENGTH: 2017
/ TYPE: DNA
/ ORGANISM: Homo sapiens
US-09-907-969-72

Query Match          9.8%; Score 507.8; DB 3; Length 2017;
Best Local Similarity 55.8%; Pred. No. 4.9e-107;
Matches 1138; Conservative 0; Mismatches 802; Indels 99; Gaps 5;

QY 175 GCGTCGATAGCAAGTAAAGTAACAGAACCATGGCTCAGTTTCCACACCTTTTGGT 234
DB 9 GCGTCGAGAGCTCGAAGAAGTCAAGGATCATGTGGCTCAGTTTCCACAGCGATGAAT 68

QY 235 GCGAGCTCGATATCTGGGCCATAACTGTAGAGGAAGAGCGAAGCATGATCAGCAGTTTC 294
DB 69 GGAGGCCAAATATGTGGGCTATTACATCTGAAGAAGCTACTAAGCATGATAACAGTTT 128

QY 295 CATAGTTTAAAGCCAATATCTGGATTCATTACTGTGTGATCAAGCTAGAAACTTTTTTTT 354
DB 129 GATAACCTCAAACCTTCAGGAGGTTACATAACAGGTGATCAAGCCGCTACTTTTCCCTA 188

QY 355 CAATCTGGGTTACCTCAACTGTTTTCACACAGATATGGGCTAGCTACATGATATAT 414
DB 189 CAGTCAGGTCCTCGGCCCGGCTTTAGCTGAATAATGGGCTTATCAGATCTGAACAAAG 248

QY 415 GATGGAAGAAATGGATCAAGTGAGTTTCCATAGCTTATGAACCTTATCAAACTGAAGCTA 474
DB 249 GATGGGAAGATGGACCAAGAGTCTCTATAGCTATGAACCTATCAAGTTAAAGTTG 308

QY 475 CAAGGATATCAGTACCTCTCTGCATCTCCCTCTGTCATGAAACAGCAACC---AGTTGCT 531
DB 309 CAGGGCCAAACAGTGGCTGTAGTCTCTCCCTCTCATCATGAACAAACCCCTATGTCTCT 368

QY 532 ATTTCTAGCGCACCATGATTTGGTATGGGAGGTATCGGCAGCATGCCACCGCTTACAGCT 591
DB 369 CCACATAATCTCTGCTCGTTTGGGATGGGAAGCATGCCCAATCTGTCTCATTCAGCCA 428

QY 592 GTTGTCTCAGTGCATATGGATC-----CATTT
DB 429 TTGCTCTCAAGTTGCACCTATAGCAACACCTTGTCTTCTGCTACTTCAGGGACCGATTT 488

QY 619 CCAGTTGTTGGAATGTCCTCAACCCCTAGTATCTTCTGTTCCACAGCAGCTGTGCCCCC 678
DB 489 CCTCCCTTAATGATGCTCTCTCCCTAGTGGCTTCTGTGTAGTACATCTCTCATTCACCAAT 548

QY 679 CTGGCTAACGGGCTCCCTCTGTATACAACTCTGCCTGCAATTTGCTCATCTCAGGCC 738
DB 549 GGAATCTGCCAGTCTCAATCAGCCTTTATCCATTCCTTATCTTCTTCAACATTTGCCCTCAT 608

QY 739 ACAATGCCCAAGAGTTCTTCTTTTAGTAGATCTGGTCCAGGGTCAAACTAAACACTAA 798
DB 609 GCATCATCTTACAGCCTGATGAGGAGGATTTGGTGTGTAGTATCCAGAGGCCAG 668

QY 799 TTACAAAGCGCAGTCATTGTGATGTGGCCAGTGTCCCAACAG-----841
DB 669 TCTCTGATTGATTAGGATCTAGTAGCTCAACTTCTCACTGCTTCCCTCTCAGGGAAC 728

QY 842 -----TGGCAGATGGGCTGTCTCTCAGTCATCAGACTGAAATACAGG 885

Db 729 TCACCTAAGACAGGGAGCTCAGAGTGGGAGTTCTCTCAGCCTTCAAGATTAAAGTATCGG 788
QY 886 CAATTTATTCAATAGTCATGACAAAACTATGAGTGGACACTTTAAACAGGTCCCAAGCAAGA 945
Db 789 CAAAAATTTAATAGTCTAGACAAAAGGCATGAGCGATACCTCTCAGGTTTTTCAAGCTAGA 848
QY 946 ACTATTCTTATGCAGTCAAGTTTACCAGGCTCAGCTGGCTTCAATATGGAATCTTTCT 1005
Db 849 AATGCCCTTCTTCAATCAATCTCTCTCAAACTCAGTAGCTACTATTGGAATCTGGCT 908
QY 1006 GACATTGATCAAGATGGAAAACTTACAGCAGAGGAATTTATCTCGCAATGCACTCATTT 1065
Db 909 GACATCGATGGTGACGGACAGTTGAAGCTGAAGAAATTTATTTCTGGCATGCACTCAT 968
QY 1066 GATGTAGCTATGTCTGGCCAAACCACTGCCACCTGTCTCTGCCCTCCAGAAATACATTCACCT 1125
Db 969 GACATGGCCAAAAGCTGGACAGCACTACCACTGACGTTTGCCTCCCGAGCTTGTCCCTCCA 1028
QY 1126 TCTTTTAGAAGAGTTGATCTGGCAGTGGTATATCTGTCTATAGCTCAACATCTGTAGAT 1185
Db 1029 TCTTTTCAAGGGGGAAAGCAAGTTGAT-----TCTGTTAATGGAACTCTGCTTTCATAT 1082
QY 1186 CAGAGGCTACCAAGAGAACCAAGTTTGAAGATGAACAACAACAAATTTAGAAAAAGAAATTA 1245
Db 1083 CAGAAAAACAAAGAAGAAGAGCT-----CAGAAAGAACTG 1118
QY 1246 CCGTTAAGCTTTGAAGATAAGAACGGGAGAACTTTTGAACGTGGCAACTGGAACTGGAG 1305
Db 1119 CCAGTTTACTTTTGGAGACAAACGGAAAGCCAACTATGAACGAGGAAACATGGAGCTGGAG 1178
QY 1306 AAACGAAAGCAAGCTCTCTCTGGAACAGCAGCGCAAGGAGCAGGAGCGCTGCCCCAGCTG 1365
Db 1179 AAGCGACCCCAAGTTTGTATGGAGCAGCAGAGGAGGAGGCTGAAACGCAAAAGCCAGAAA 1238
QY 1366 GAGCGCGGAGCAGGAGGAAGAGCGTGTAGCGCCAGGAGCAAGAGCGCAAAAAGACAA 1425
Db 1239 GAGAAGGAAGAGTGGAGCGGAAACAGAGAGAACTGCAAGAGCAAGAAATGGAAGAAGCAG 1298
QY 1426 CTGGAACCTGGAGAACAACTCGAAAGCAGCGGGAGCTTAGAACGCGCAGAGAGAGGAGAG 1485
Db 1299 CTGGAGTTGGAGAAACCGCTTGGAGAAAACAGAGAGAGCTGGAGAGACAGCGGAGGAAGAG 1358
QY 1486 AGGAGAAAGAAATTTGAGAGGCGAGAGCTGCAAAACGGGAACCTTGAAGGCAACGACAA 1545
Db 1359 AGGAGAAAGGAGATAGAAAGACGAGAGCGCAAAAACAGGAGCTTGGAGAGACAGCCCT 1418
QY 1546 CTTGAGTGGGAACGGAATCGAAGGCAAGAACTACTATAATCAAAAGAAAACAAAGACAG 1605
Db 1419 TTAGAAATGGGAAGAAGACTCGCTCGGACAGGCTGTCTCAGTCAAGAACAGGGAACAAGAA 1478
QY 1606 GACATAGTTGTACTGAAAGCAAGAAAGAAAGACTTTTGGAAATTTGAATTTAGAAGCTCTAAT 1665
Db 1479 GACATTTGTCAAGCTCAGCTCCAGAAAGAAAAGTCTCCACCTGGAACTTGAAGCAGTGAAT 1538
QY 1666 GATAAAAAGCATCACTAGAAAGGGAACCTTCAAGATATCAGATGTCGATTGACCCCAA 1725
Db 1539 GGAAAAACATCAGCAGATCTCAGGCGAGACTACAAAGATGTCCAAATCAGAAAGCAACAA 1598
QY 1726 AGGCAAGAAATTTAGAGCACAACAAATCTTAGAGAGTTTGAGAAATTCGCGAAATCACCCAT 1785
Db 1599 AAGACTGAGCTAGAGTTTGGATAAAACAGTGTGACCTGGAAATTTATGGAATCAACAA 1658
QY 1786 CTACAGCAACAAATTAACAGAAATCTCAGCAAAATGCTTTGGAAGAGACTTTATCCAGAAAAACAG 1845
Db 1659 CTTCAACAAGAGCTTTAAGGAATATCAAAATAAGCTTATCTATCTGCTCCCTGAGAAGCAG 1718
QY 1846 ATACTCAATGACCAATTAACAAAGTTTCAGCAGACAGCTTTTGCACAGAGATTCACCTGTT 1905
Db 1719 CTTATTAACGAAAGAAATTAACAAACATGCAGCTCAGTAACACACCTGATTCAGGATCAGT 1778
QY 1906 ACACCTTAAAGAGCCCTTAGAAGCAAAAGAACTTAGCTCGGCGAGCACCTACGAGACCAACTG 1965
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| | | | |
|--|------|---|------|
| Db | 1779 | TTACTTCATAAAAGTCATCAGAAAAGGAAGAAATATATGCGAAAGACTTTAAAGACAAATTA | 1838 |
| Qy | 1966 | GATGAAGTGGAGAAAGAACTAGATCAAAAGCTACAGGAGATTGATATTTTCAATAATCAG | 2025 |
| Db | 1839 | GATGCTCTTTGAAAAAGAAAGTGCATCTAAGCTCTCAGAAATGGAATTCATTTAAACAATCAG | 1898 |
| Qy | 2026 | CTGAAGGAATTAAGAGAAATACACATAAGCAACAACTCCAGAAAGCAAAAGTCCATGGAG | 2085 |
| Db | 1899 | CTGAAGGAATCTCAGAGAAAGCTATATAACACAGCAGTTAGCCCTTGAACAATCTTCATATA | 1958 |
| Qy | 2086 | GCTGAACGACTGAACACAGAAAGAAACAAAGAACGAAAGATCATAGAAATTAGAAAAACAAA | 2144 |
| Db | 1959 | ATCAAAAGCTGACAAATTTGAAGGAATCGAAGGAAGAAAGATTAGAGCAAAAGAAAAAAA | 2017 |
| RESULT 12 | | | |
| US-09-827-271-72 | | | |
| ; Sequence 72, Application US/09827271 | | | |
| ; Publication No. US20030165504A1 | | | |
| ; GENERAL INFORMATION: | | | |
| ; APPLICANT: Retter, Marc W. | | | |
| ; APPLICANT: Fanger, Gary R. | | | |
| ; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND | | | |
| ; TITLE OF INVENTION: DIAGNOSIS OF OVARIAN CANCER | | | |
| ; FILE REFERENCE: 210121.462C6 | | | |
| ; CURRENT APPLICATION NUMBER: US/09/827, 271 | | | |
| ; CURRENT FILING DATE: 2001-04-04 | | | |
| ; NUMBER OF SEQ ID NOS: 461 | | | |
| ; SOFTWARE: PastSeq for Windows Version 3.0 | | | |
| ; SEQ ID NO 72 | | | |
| ; LENGTH: 2017 | | | |
| ; TYPE: DNA | | | |
| ; ORGANISM: Homo sapien | | | |
| US-09-827-271-72 | | | |
| Query Match 9.88; Score 507.8; DB 3; Length 2017; | | | |
| Best Local Similarity 55.88; Pred. No. 4.9e-107; | | | |
| Matches 1138; Conservative 0; Mismatches 802; Indels 99; Gaps 5; | | | |
| Qy | 175 | GGCGTCGATTTAGCAAGGTAAAGTAAACAGAACCATGGCTCAGTTTCCACACCTTTTGGT | 234 |
| Db | 9 | GGCTGAGAGCTGCAAGAAGAGTCAGGATCATGATGCTCAGTTTCCACAGCGATGAAT | 68 |
| Qy | 235 | GGCAGCGCTGATATCTGGGCGATAA CTGTAGAGGAAGAGCGAAGCATGATCAGCAGTTC | 294 |
| Db | 69 | GGAGGCGCAATATGTGGGCTATTACATCTGAAGAACGTACTAAGCATGATAAACAGTTT | 128 |
| Qy | 295 | CATAGTTTAAAGCCAATATCTGGATTCATTCTGTTGATCAAGCTAGAACTTTT | 354 |
| Db | 129 | GATAAACCCTCAACCTTCAGGAGGTACATAACAGGTATCAAGCCCGTACTTTTTCCTA | 188 |
| Qy | 355 | CAATCTGGGTACCTCAACCTGTTTATAGCACAGATATGGGCATCTAGCTGACATGAATAT | 414 |
| Db | 189 | CAGTCAGTCTGCGGCGCCGGTTTATAGCTGAATATATGGGCTTATCAGATCTGAACAAG | 248 |
| Qy | 415 | GATGGAGAAATGGATCAAGTGGAGTTTCCATAGCTATGAACCTTATCAAACTGAAGCTA | 474 |
| Db | 249 | GATGGGAAGATGGACCAAGAGTTCTCTATAGCTATGAACCTATCAAGTTTAAAGTTG | 308 |
| Qy | 475 | CAAGGATATCAGCTACCTCTGCACTTCCCGCTGTCATGAACACAGCAACC---AGTTGCT | 531 |
| Db | 309 | CAGGGCCAACAGCTGCTGTAGTCTCCCTCTCTATATGAACAAACCCCTATGTTCTCT | 368 |
| Qy | 532 | ATTTCTAGCGCACCAAGATTTGGTATGGAGGTATCGCCAGCATGCCACCGCTTACAGCT | 591 |
| Db | 369 | CCACTAATCTCTGCTCGTTTGGTGGAGAGCATGCCAATCTGTCCATTTCATCAGCCA | 428 |
| Qy | 592 | GTTGCTCCAGTGCCTAATGGGATC-----CATT | 618 |
| Db | 429 | TTGCTCCAGTTGCACCTTATAGCAACACCCCTTGTCTCTGCTACTTTCAGGGACCAAGTAT | 488 |
| Qy | 619 | CCAGTTGTTGAATGTCTCCAACTAGTATCTTCTGTTCCACAGCAGCTGTGCCCCC | 678 |

| | | | |
|----|------|--|------|
| Db | 489 | CCTCCCTAAATGATGCTGCTCCCTCTAGTGCCTCTCTGTTAGTACATCTCATTTACCAAT | 548 |
| Qy | 679 | CTGGCTTAGGGGGCTCCCTCTGTATACAACTCTGCTGCAATTTGCTCATCTGTCAGGC | 738 |
| Db | 549 | GGAACTGCCAGTCTCATTCAGGCTTTATCCATTCCTTATCTTCTTCAACATTCGCTCAT | 608 |
| Qy | 739 | ACATTTGCCAAAGAGTTCTTCTTTTAGTAGTATGCTGCTCCAGGGTCAAACTAAACACTAAA | 798 |
| Db | 609 | GCATCATCTTACAGCTGATGATGGAGATTTGGTGGTGTAGTAGTATCCAGAAAGGCCAG | 668 |
| Qy | 799 | TTACAAAGGACAGTCATTTGATGTGGCAGTGTCCACCCAG-----841 | |
| Db | 669 | TCTCTGATTTAGTATCTAGTCAACTTCTCACTGCTTCTCCCTCTCAGGGAAC | 728 |
| Qy | 842 | -----TGGCAGAGTGGGCTGTTCTCTCAGTCATCAAGACTGAAATACAGG | 885 |
| Db | 729 | TCACCTAAGACAGGGACCTCAGATGGGCAGTTCTCAGCCTTCAAGATTTAAAGTATCGG | 788 |
| Qy | 886 | CAATTATTCAATAGTATGACAAACTATGAGTGGACACTTAAACAGGTCCCCAAGCAGA | 945 |
| Db | 789 | CAAAAATTTAATAGTCTAGACAAAGGCATGAGCGGATACCTCTCAGGTTTTCAGCTAGA | 848 |
| Qy | 946 | ACTATTCTTATGAGTCAAGTTTACCAAGGCTCAGCTGGCTTCAATATGGAATCTTTCT | 1005 |
| Db | 849 | AATGCCCTTCTTTCAGTCAAACTCTCTCAACTCAGCTAGTACTATTTTGGACTCTGGCT | 908 |
| Qy | 1006 | GACATTTGATCAGATGAAACCTTACAGCAGAGGAATTTATCTGGCAATGCACTCATTT | 1065 |
| Db | 909 | GACATCGATGGTACCGGACAGTTGAAAGCTGAAGATTTATTTCTGGGGATGCACTCACT | 968 |
| Qy | 1066 | GATGTAGTATGCTGGCCAAACCACTGCCACCTGCTCTGCTCCAGAAATACATTTCACT | 1125 |
| Db | 969 | GACATGCCCAAGCTGACAGCCACTACCACTGACGTGGCTCCCGAGCTTGTCCCTCCA | 1028 |
| Qy | 1126 | TCTTTTAGAAGATTGATCTGCGAGTGTATATCTGTCTATAGCTCAACATCTGTAGAT | 1185 |
| Db | 1029 | TCTTTCAGAGGGGGAAGCAAGTTGAT-----TCTGTTAATGGAACCTGCGCTTCATAT | 1082 |
| Qy | 1186 | CAGAGCTACACAGAGAACCCAGTTTGTAGAGATGMAACAACAACAAATTTAGAAAAAATTA | 1245 |
| Db | 1083 | CAGAAAACAAGAAAGAGCGCT-----CAGNAGAAATCG | 1118 |
| Qy | 1246 | CCTGTAACTGTTGAAGATAAGAACCGGAGAACTTTGAACGTGGCAACCTTGGAACTGGAG | 1305 |
| Db | 1119 | CCAGTTACTTTTTCAGGACAAACCGAAAGCCAACTATGAACGAGGAAACATGGAGCTGG | 1178 |
| Qy | 1306 | AAACGAAGGCAAGCTCTCTGGAAACAGCAGCCGACGAGGAGGAGCGCTGGCCACGCTG | 1365 |
| Db | 1179 | AAGCGACGCCAAGTGTGTATGAGCAGCAGCAGGAGGAGGCTGAACCGAAAGCCAGAAA | 1238 |
| Qy | 1366 | GAGCGGCGGACGAGGAGAGGAGGAGCGGTGAGCGCCAGGAGCAAGAGCGCAAAAGACAA | 1425 |
| Db | 1239 | GAGAAGGAAGTGGGAGCGGAAACAGAGAGAACTGCAAGGCAAGATTTGAAGAGCAG | 1298 |
| Qy | 1426 | CTGGAACTGAGAGAACCACTGGAAAGACAGCGGAGCTAGAAACGGCAGAGAGAGGAGAG | 1485 |
| Db | 1299 | CTGGAGTTGGAGAAACGCTTTGGAGAAACAGAGAGAGCTGAGAGACAGCGGAGGAGAG | 1358 |
| Qy | 1486 | AGGAGGAAAGAAATTTAGAGCGGAGAGGCTGCAAAACGGGAACTTTGAAGGCAACGACAA | 1545 |
| Db | 1359 | AGGAGAAAGAGATAGAAAGACGAGGCGCAAAAACAGGAGCTTTAGAGACAACCGCGT | 1418 |
| Qy | 1546 | CTTGAGTGGGAACGGAAATCGAAGCAAGAACTACTAAATCAAGAAACAAAGAAACAAGAG | 1605 |
| Db | 1419 | TTAGATGGGAAAGAGCTCCGTGCGCAGAGAGCTGCTCAGTCAGAGACCGAGGAAACAGAA | 1478 |
| Qy | 1606 | GACATAGTTGTACTGAAAGCAAAAGAAAGCTTTTGGAAATTTGAATTTGAAGCTCTAAAT | 1665 |
| Db | 1479 | GACATTTCTCAGCTGAGCTCCAGAAAGAAAGTCTCCACCTGGAACTGGAAGAGCTGAAT | 1538 |
| Qy | 1666 | GATAAAAGCATCAACTAGAAAGGAAACCTTCAAGATATCAGATGTCTGATTCAGCACCAAA | 1725 |
| Db | 1539 | GGAAAAATCAGCAGATCTCAGGCGAGACTACAAGATGTCCAAATTCAGAAAGCAACAA | 1598 |

1299 CTGGAGTTGGAGAAACGCTTGAGAGAAACAGAGAGAGCTGGAGAGACAGCGGAGGAGAG 1358
1486 AGGAGGAAAGAAATTTAGAGGCGAGAGGCTGCAAAACGGGAACTTGAAGGCAACGACAA 1545
1359 AGGAGAAAGAGATAGAAAGACGAGAGGCGCAAAACAGGAGCTTGAGAGACAAACGCCGT 1418
1546 CTTGAGTGGGACCGGATCGAAGGCGAGAGGCTTAAATCAAGAAACAAAGCAAGAG 1605
1419 TTAGAATGGGAAGACTCCGTCGGCAGGAGCTGCTAGTCAGAGACGAGGAGCAAGAA 1478
1606 GACATAGTTGTACTGAAAGCAAGAAAGAAAGACTTTGGAAATTTGAATTTAGAAGCTCTAAAT 1665
1479 GACATTTGTCAGGCTGAGCTCCAGAAAGAAAGTCTCCACTGGAAGTGAAGCAGTGAAT 1538
1666 GATAAAAGCATCAACTAGAGGGAATTTCAAGATATCAGATGTCGATTTGACCAACCA 1725
1539 GGAACACATCAGCAGATCTCAGGACAGCTACAGAGATGTCGAATCAGAAAGCAACACAA 1598
1726 AGGCAAGAAATTCAGAGCACAACAAATCTAGAGAGTTGAGAAATGCGCGAAATCAACAT 1785
1599 AAGACTGAGCTAGAGTTTGGATAAACACAGTGTGACCTGGAATTTATGGAATTCAAACAA 1658
1786 CTACAGCAAAATTCAGAGAACTCTCAGCAAAATGCTTGAAGAGCTTATTCAGAGAAACAG 1845
1659 CTTCAACAGAGCTTAAAGATATCAAAATAGCTTATCTATCTGCTCCCTGAGAGCAG 1718
1846 ATACTCAATGACCAATTTAAACAAAGTTTCAGCAGAAACAGTTTGCACAGAGATTCATTGTT 1905
1719 CTATTAAACGAAAGATTTAAACACATGCAGCTCAGTAAACACACCTGATTCAGGGATCAGT 1778
1906 ACATTTAAAGAGCTTAGAGCAAAAGAAAGTCTGCGCAGCAGCTACGAGACCAACTG 1965
1779 TTACTTCATATAAAGTCATCAGAAAGGAAAGTATGCGCAAGAGCTTAAAGCAATTA 1838
1966 GATGAAGTGAGAGAAAGAACTAGATCAAAACTACAGGAGATGATATTTTCAATAATCAG 2025
1839 GATGCTCTTGAAGAAAGAACTGATCTAGCTCTCAGAAATGATTCATTTAAACATCAG 1898
2026 CTGAAGGAACTAGAGAAATACAAATAAGCAACAACTCCAGAGCAAAAGTCCATGGAG 2085
1899 CTGAAGGAACTCAGAGAAAGCTAATAACACAGCAGTTAGCCCTTGAACAACTTCATAAA 1958
2086 GCTGAAGCAGTGAACAGAGAAACAGACGAGAAAGTATCAGAAATTTAGAAACAAAA 2144
1959 ATCAACGTCGACAAATTTGAAGGAAATCGAAAGAAAGATTTAGAGCAAAAGAAAAAAA 2017

RESULT 14
US-10-860-790-72
; Sequence 72, Application US/10860790
; Publication No. US20050031634A1
; GENERAL INFORMATION:
; APPLICANT: Bangur, Chaitanya S.
; APPLICANT: Retter, Marc W.
; APPLICANT: Fanger, Gary R.
; APPLICANT: Hall, Paul
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
; TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
; FILE REFERENCE: 210121.462C11
; CURRENT APPLICATION NUMBER: US/10/860,790
; CURRENT FILING DATE: 2004-06-02
; NUMBER OF SEQ ID NOS: 624
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 72
; LENGTH: 2017
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-860-790-72

Query Match 9.8%; Score 507.8; DB 8; Length 2017;
Best Local Similarity 55.8%; Pred. No. 4.9e-107;
Matches 1138; Conservative 0; Mismatches 802; Indels 99; Gaps 5;

175 GCGCTCGATTAGCAAGGTAAAGTAAACAGAACCATGGCTCAGTTTCCAAACACCTTTTGGT 234
9 GCTCAGAGCTGCAAGAGAGAGTCAAGATCATGATGGCTCAGTTTCCCAACAGCATGAAT 68
235 GCGACCTGGATATCTGGGCCATTAATCTGTAGAGGAAAGAGCGAAGCATGATCAGCAGTTTC 294
69 GGAGGCCAAATATCTGGGCTATTACATCTGAAGAACGTACTAAGCATGATTAACACAGTTT 128
295 CATAGTTTAAAGCCAAATATCTGGATTCAATCTGCTGATCAAGCTAGAGAACTTTTTTTTTT 354
129 GATAACCTCAAACTTTAGGAGTTACATACAGGTGATCAAGCCGCTATTTTTCCTA 188
355 CAATCTGGGTTACCTCAACCTTTTACACAGATATGGCAGCTAGCTGACATGATTAATAT 414
189 CAGTCAGGCTGCGCGGCCCGGTTTTAGCTGAAATATGGCCTTATCAGATCTGAACAAG 248
415 GATGAAGAAATGATCAAGTGGAGTTTCCATAGCTATGAAACTTATCAAACTGAAGCTA 474
249 GATGGAAAGATGGACCAAGAGTTCTATAGCTATGAAACTCATCAAGTTTAAAGTTG 308
475 CAAGGATATCAGTACCTCTGCACTTCCCTCTGTCATGAAACAGCAACC---AGTTGCT 531
309 CAGGGCAACAGCTGCTGCTGCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCTCT 368
532 ATTTCTAGCGCACCAAGCAATTTGGTATGGAGGATATGGCAGCATGCCACCGCTTACAGCT 591
369 CCATTAATCTCTGCTGCTTTTGGGATGGGAGCATGCCCAATCTGTCCATTCATCAGCCA 428
592 GTTGCTCCAGTGGCAATGGGATC-----CAAT 618
429 TTGCTCTCCAGTTGCACCTATAGCAACACCTTTGCTCTTCTGCTACTTTCAGGGACCATGAT 488
619 CCAGTTGTTGGAATGTCCTCAACCTTAGTATCTCTGTTCCCAACAGCAGCTGTGCCCCCT 678
489 CTTCCCTTAATGATGCTGCTCTCCCTAGTGCTCTCTGTTAGTACCTCTCATTACCAGAT 548
679 CTGGCTAAACGGGCT 738
549 GGAATGCTCCAGTCTCATTTCAGCTTTATCCATTCCTTCTCTCAACATTTGCTCAT 608
739 ACATGCGCAAGAGTTCT 798
609 GCATCATCTTACAGCTGATGATGGAGGATTTGGTGGTGTAGTATCCAGAGGCGCCAG 668
799 TTACAAAGGCGACAGTCATTGATGGCGAGTGTCCCAACAG----- 841
669 TCTCTGATTTAGGATCTAGTAGTCAATCTCTCACTGCTTCTCTCTCTCTCTCTCTCT 728
842 -----TGGCAGAGTGGGCTGTTCTCTCAGTCATCAAGACTGAAATACAGG 885
729 TCACCTAAGACAGGGACCTCAGAGTGGGAGTTCTCTCAGCTTCAAGATTTAAAGTATCGG 788
886 CAATTAATCAATAGTATGACAAACTATGAGTGGACACTTAACAGGTCCCAAGCAAGA 945
789 CAAAAATTTAATAGTCTAGACAAAGGATGAGCGGATACCTCTCAGGTTTTCAGGCTAGA 848
946 ACTATTCTTATGAGTCAAGTTTACCAAGGCTCAGTGGCTTCAATATGGAATCTTTCT 1005
849 AATGCCCTTCTCTCAGTCAAACTCTCTCAACATCAGTAGTACTATTTGGACTCTGGCT 908
1006 GACATTGATCAAGATGGAAACTTACAGCAGAGGAAATTTATCTGGCAATGCACTCAT 1065
909 GACATCGATGGTGACGAGAGTTGAAAGCTGAAAGATTTATCTGGCGATGCACCTCACT 968
1066 GATGTAGTATGCTGGCCAAACCACTGCCACCTGCTCTGCTCTCAGNATACATTTCCACCT 1125
969 GACATGGCCAAAGCTGGACAGCCACTACCACTGACGTTGCTCTCCGAGCTTGTCCCTCCA 1028
1126 TCTTTTAGAGAGTTCGATCTGCGAGTGGTATATCTGTATAAGCTCAACATCTGTAGAT 1185
1029 TCTTTTAGAGAGGAGGAAAGCAAGTTGAT-----TCTGTTAATGNACTCTGCTCTCATAT 1082
1186 CAGAGGCTACAGAGGAACCAAGTTTATAGAGATGAACAAACAAATTTAGAAAAGAAATTA 1245

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Db 1083 CAGAAAACAAGAAAGAGCGCT-----CAGAAAGAACTG 1118
Qy 1246 CTGTAAACGTTTGAAGATAAGACGGGAGAACTTTGAACGTGGCAACCTGGAACCTGGAG 1305
Db 1119 CCAGTTACTTTTGAAGACAACCGAAAGCCACTATGAACGAGGAAACATGGAGCTGGAG 1178
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RESULT 15

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US-09-764-881-55
; Sequence 55, Application US/09764881
; Publication No. US20020086821A1
; GENERAL INFORMATION:
; APPLICANT: Rosen et al.
; TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
; FILE REFERENCE: PT207
; CURRENT APPLICATION NUMBER: US/09/764,881
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; CURRENT FILING DATE: 2001-01-17
; Prior application data removed - refer to PALM or file wrapper
; NUMBER OF SEQ ID NOS: 192
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 55
; LENGTH: 568
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: SITE
; LOCATION: (481)
; OTHER INFORMATION: n equals a,t,g, or c
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; LOCATION: (562)
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US-09-764-881-55
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Query Match          9.7%; Score 503.6; DB 3: Length 568;
Best Local Similarity 96.9%; Pred. No. 2.1e-106;
Matches 554; Conservative 0; Mismatches 13; Indels 5; Gaps 4;

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Search completed: February 14, 2006, 06:40:01
Job time : 3793 secs

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| 1 | 551.6 | 10.6 | 5813 | 3 | US-09-949-016-1739 | Sequence 1739, Ap |
| 2 | 507.8 | 9.8 | 2017 | 3 | US-09-404-879A-72 | Sequence 72, Appl |
| 3 | 507.8 | 9.8 | 2017 | 3 | US-09-338-933-72 | Sequence 72, Appl |
| 4 | 507.8 | 9.8 | 2017 | 3 | US-09-215-681-72 | Sequence 72, Appl |
| 5 | 507.8 | 9.8 | 2017 | 3 | US-09-216-003A-72 | Sequence 72, Appl |
| 6 | 507.8 | 9.8 | 2017 | 3 | US-09-667-857-72 | Sequence 72, Appl |
| 7 | 507.8 | 9.8 | 2017 | 3 | US-10-198-053-72 | Sequence 72, Appl |
| 8 | 507.8 | 9.8 | 2017 | 3 | US-09-827-271-72 | Sequence 72, Appl |
| 9 | 174 | 3.3 | 174 | 3 | US-09-513-999C-27927 | Sequence 27927, A |
| 10 | 165.2 | 3.2 | 2873 | 3 | US-08-630-915A-193 | Sequence 193, App |
| 11 | 165.2 | 3.2 | 2873 | 3 | US-09-879-957-193 | Sequence 193, App |
| 12 | 163.8 | 3.2 | 747 | 3 | US-08-630-915A-39 | Sequence 39, Appl |
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| 14 | 153.4 | 3.0 | 531 | 3 | US-08-404-879A-5 | Sequence 5, Appl |
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| 19 | 153.4 | 3.0 | 531 | 3 | US-10-198-053-5 | Sequence 5, Appl |
| 20 | 153.4 | 3.0 | 531 | 3 | US-09-827-271-5 | Sequence 5, Appl |
| 21 | 149.2 | 2.9 | 480 | 3 | US-09-404-879A-60 | Sequence 60, Appl |
| 22 | 149.2 | 2.9 | 480 | 3 | US-09-338-933-60 | Sequence 60, Appl |
| 23 | 149.2 | 2.9 | 480 | 3 | US-09-215-681-60 | Sequence 60, Appl |
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RESULT 3
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; Sequence 72, Application US/09338933
; Patent No. 6488931
; GENERAL INFORMATION:
; APPLICANT: Mitcham, Jennifer Lynn
; APPLICANT: King, Gordon E.
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY OF
; TITLE OF INVENTION: OVARIAN CANCER
; FILE REFERENCE: 210121.462C1
; CURRENT APPLICATION NUMBER: US/09/338,933
; CURRENT FILING DATE: 1999-06-23
; NUMBER OF SEQ ID NOS: 312
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 72
; LENGTH: 2017
; TYPE: DNA
; ORGANISM: Homo sapien
; US-09-338-933-72

Query Match 9.8%; Score 507.8; DB 3; Length 2017;
Best Local Similarity 55.8%; Pred. No. 7.3e-111;
Matches 1138; Conservative 0; Mismatches 802; Indels 99; Gaps 5;

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| Db | 609 | GCATCATCTTACAGCCTGATGATGGAGGATTTGGTGGTGTAGTAGTCCAGAAGGCCAG | 668 |
| Qy | 799 | TTACAAAGGCACAGTCAATTTGATGTGGCAGTGTCCACACAG----- | 841 |
| Db | 669 | TCTCTGATTGATTTAGGATCTTAGTAGCTCAACTTCTCTCAATGTCTTCCCTCTCAGGGAAC | 728 |
| Qy | 842 | -----TGGCAGAGTGGGTGTTCCTCAGTCACTCAAGACTGAAATACAG | 885 |
| Db | 729 | TCACCTAAGACAGGGACCTCAGGTGGCAGTTCTCAGCCTTCAAGATTTAAAGTATCG | 788 |
| Qy | 886 | CAATTATTCAATAGTCATGACAAACTATAGTGGGACATTAAACAGGTCCCAAGCAAGA | 945 |
| Db | 789 | CAAAATTTTAATAGTCTAGACAAAGGCATGAGCGGATACCTCTCAGGTTTCAAGCTAGA | 848 |
| Qy | 946 | ACTATTCTTATGCAAGTATTACACAGGCTCAGCTGGCTTCATATGGAATCTTTCT | 1005 |
| Db | 849 | AATGCCCTTCTTCAGTCAATCTCTCTCAAACTCAGCTAGCTACTATTTTGGACTCTGGCT | 908 |
| Qy | 1006 | GACATTGATCAAGATGGAACCTTACAGACAGAGAAATTTATCTGCGCAATGCACTCAT | 1065 |
| Db | 909 | GACATCGATGGTGACGACAGTTGAAAGCTGAAGATTTATTCTGGCGATGCACTCACT | 968 |
| Qy | 1066 | GATGTAGCTATGTCGGCCAAACACTGCCACCTGTCTGCTCCAGAAATACATTCACCT | 1125 |
| Db | 969 | GACATGCCAAAGCTGGACAGCCACTACCACTGACGTTGCTCCCGAGCTGTGCTCCCA | 1028 |
| Qy | 1126 | TCCTTTTGAAGAGTTTCGATCTGCGAGTGGTATATCTGTCTATAAGCTCAACATCTGTAGAT | 1185 |
| Db | 1029 | TCCTTTCAGAGGGGGAAGCAAGTTGAT-----TCGTTAATGGAACCTGTCCTTTCATAT | 1082 |
| Qy | 1186 | CAGAGGCTACGAGGAACCAAGTTTGAAGATGAAACAAACAAATTTAGAAAAGAAATTA | 1245 |
| Db | 1083 | CAGAAAACAAGAGAGAGCCT-----CAGAGAAACTG | 1118 |
| Qy | 1246 | CCTGTAACGTTTGAAGATAAGAGCGGAGAACTTTGAACGTGGCACTCTGGAACTGGAG | 1305 |
| Db | 1119 | CCAGTTACTTTTCAGGACAAACCGAAAGCCAACTATGAACGAGAAACATGAGCTGGAG | 1178 |
| Qy | 1306 | AAACGAAGGCAAGCTCTCTGGAAACAGCAGCGCAAGGAGCAGGCGCTTGGCCAGCTG | 1365 |
| Db | 1179 | AAGCGACGCCAAGTGTGTATGGAGCAGCAGAGGGAGGCTGAACCAAGCCCAAGAA | 1238 |
| Qy | 1366 | GAGCGGCGAGCAGGAGAGAGAGGAGCGGTGAGCGCCAGGAGCAAGAGCGCAAAAGACAA | 1425 |
| Db | 1239 | GAGAGGAAGATGGGAGCGGAACAAGAGAGAACTGCAAGCAGAGATGGAAGAGCAG | 1298 |
| Qy | 1426 | CTGGAACTGGAGAGCAACTGGAAAGACGCGGAGCTTAGAAACGCGAGAGAGGAGGAG | 1485 |
| Db | 1299 | CTGGAGTTGGAGAAACGCTTTGGAGAAACAGAGAGAGCTGGAGACACAGCGGAGGAAGAG | 1358 |
| Qy | 1486 | AGGAGGAAGAAATTTAGAGCGCAGAGGCTGCAAAACGGGAACTTGAAGGCAACGACAA | 1545 |
| Db | 1359 | AGGAGAAAGGAGATAGAAAGACGAGAGGCGAGCAAAACAGGAGCTTTGAGACACACGCCGT | 1418 |
| Qy | 1546 | CTTGAGTGGGAACGGAATCGAAGGCAAGAACTACTAAATCAAGAAAACAAGAACAGAG | 1605 |
| Db | 1419 | TTAGATGGGAAGAGCTCCGTGCGCAGAGCTGCTCAGTCAGAGACACAGGAAACAGAA | 1478 |
| Qy | 1606 | GACATAGTTGTACTGAAAGCAAAAGAACTTTTGAAATTTGAAATTTAGAGCTCTTAAAT | 1665 |

RESULT 4

```

RES001.4
US-09-215-681-72
; Sequence 72, Application US/09215681A
; Patent No. 6528253
; GENERAL INFORMATION:
; APPLICANT: Mitcham, Jennifer L.
; APPLICANT: Frudakis, Tony N.
; APPLICANT: King, Gordon E.
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSIS
; TITLE OF INVENTION: OF OVARIAN CANCER
; FILE REFERENCE: 210121.463
; CURRENT APPLICATION NUMBER: US/09/215,681A
; CURRENT FILING DATE: 1998-12-17
; NUMBER OF SEQ ID NOS: 310
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 72
; LENGTH: 2017
; TYPE: DNA
; ORGANISM: Homo sapien
; US-09-215-681-72

```

| Query Match | 9.8% | Score 507.8; | DB 3; | Length 2017; |
|----------------------------|-------|---------------------|------------|--------------|
| Best Local Similarity | 55.8% | Pred. No. 7.3e-111; | | |
| Matches 1138; Conservative | 0; | Mismatches 802; | Indels 99; | Gaps 5 |

| | | | |
|----|-----|--|-----|
| Qy | 175 | GGCGTGGATTGACAAAGGTAAAGGTAAACAGAACCAATGGCTCAGTTTCCACACACCTTTTGGT | 234 |
| | | | |
| Db | 9 | GGCTGAGAGCTGCAAGAAGAAGTCAGGATCATGTGCTCAGTTTCCACAGCGATGAAT | 68 |
| | | | |
| Qy | 235 | GGCAGCCTGGATATCTGGGCCATAATCTGAGAGGAAGAGCGAAGCAGTATGATCAGCAGTTTC | 294 |
| | | | |
| Db | 69 | GGAGGGGCCAATAATGTGGGCTATTACATCTGAGAAACGTACTAAGCATGATTAACAGTTT | 128 |
| | | | |
| Qy | 295 | CATAGTTTAAAGCCCAATATCTGGATTTCATTACTGGTGATCAAGCTAGAAACTTTTTTTTTT | 354 |
| | | | |
| Db | 129 | GATTAACCTCAACCTTCAGGNGTTACATAACAGGTGATCAAGCCCGTACITTTTTTTCCTA | 188 |
| | | | |
| Qy | 355 | CAATCTGGGTGTAACCTCAACCTGTTTTAGCACAGATATGGGCACATAGCTGCATGATGAATAT | 414 |

Db 189 CAGTCAGGCTGCGCGCCCGGTTTTAGCTGAAATATGGCCCTTATCAGATCTGAACAAG 248
QY 415 GATGGAAGAAATGATCAAGTGGAGTTTTCATAGCTATGAACCTTATCAAACTGAAGCTA 474
Db 249 GATGGAAGATGGACAGCAAGAGTTCTCTATAGCTATGAACCTATCAAGTTAAAGTTG 308
QY 475 CAAGGATATCAGCTACCTCTGCACTTCCCTCTGTCATGAACAGCAACCC---AGTTGCT 531
Db 309 CAGGCGCAACAGCTGCTGTAGTCTCCCTCTCTATCATGAACAACCCCTATGTTCTCT 368
QY 532 ATTTCTAGCCACCACATTTGATGAGGAGTATCGCCAGCATGCCCGCTTACAGCT 591
Db 369 CCACCTAATCTCTGCTCGTTTGGGATGGGAAGCATGCCCAATCTGTCCATTCATCAGCCA 428
QY 592 GTTGCTCCAGTCCCAATGGGATC-----CAAT 618
Db 429 TTGCTCTCAGTTGCACTATAGCAACCCCTTGTCTTCTGCTACTTCAGGGACCAGTAT 488
QY 619 CCAGTTGTTGGAATGCTCCAAACCCTAGTATCTTCTGTTTCCCAAGCAGCTGTGCCCCC 678
Db 489 CTTCCCTAATGATGCTGCTCCCTAGTGCCTTCTGTTAGTACATCTCTCAATACCAAT 548
QY 679 CTGGCTAAAGGGGCTCCCTCTGTTATACAACTCTGCCCTGCAATGCTCATCTGCAGCC 738
Db 549 GGAAGTCCAGTCTCATCAGCCTTTATCCATTCCTTATCTTCTTCAACATTCGCTCAT 608
QY 739 ACATGTCCAAAAGTTCTTCTTTAGTAGATCTGGTCCAGGGTCAAACTAAACACATAA 798
Db 609 GCATCATCTTACGCTGATGAGGAGGATTTGGTGGTGTAGTAGTATCAGAAAGGCCAG 668
QY 799 TTACAAAGGCAAGTCATTTGATGTGGCCAGTGTCCCAACAG----- 841
Db 669 TCTCTGATTGATTAGGATCTAGTAGCTCAATCTCCTCACTGCTTCCCTCTCAGGGAAC 728
QY 842 -----TGCAGAGTGGCTGTCTCAGTCATCAGACTGGAATACAGG 885
Db 729 TCACCTTAAGACAGGGACCTCAGAGTGGCGAGTTCTCCAGCCTTCAAGATTAAAGTATCGG 788
QY 886 CAATATTCAATAGTCATGACAAAACATATGAGTGGACACTTAACAGGTCCCAAGCAAGA 945
Db 789 CAATAATTTAATAGTCTAGCAAAAGGCATGAGCGGATACCTCTCAGTCTTCAAGCTAGA 848
QY 946 ACTATTCCTTATGAGTCAAGTTTACACAGGCTCAGCTGGCTTCAATATGGAATCTTCT 1005
Db 849 AATGCCCTTCTTCAGTCAAACTCTCTCAAACTCAGCTAGCTACTATTTGGACTCTGGCT 908
QY 1006 GACNTTGATCAAGATGGAACCTTACAGCAGAGGATTTATCTGGCAATGCACCTCAT 1065
Db 909 GACATCGATGGTGACGGACAGTTGAAAGCTGAAGAATTTATTTCTGGCGATGCACCTCACT 968
QY 1066 GATGTAGCTATGTCTGGCAACACCTGCGCACTGTCTCTGCCCTCCAGAATACATTCACCT 1125
Db 969 GACATGGCCAAAGCTGGAACAGCACTACCACTGACGTTCCTCCGAGCTTGTCCCTCCA 1028
QY 1126 TCTTTTGAAGAGTTCGATCTGCGAGTGGTATATCTGTATAGCTCAACATCTGTPAGAT 1185
Db 1029 TCTTTTCAAGGGGGAAGCAAGTTGAT-----TCTGTAAATGGAACCTCTGCTTCTCATAT 1082
QY 1186 CAGAGCTACCAGAGAACCTGTTTGAAGATGAACAACAATTAGAAGAAATTA 1245
Db 1083 CAGAAACACAAGAAGAGAGCT-----CAGAAGAAACTG 1118
QY 1246 CTGTAACTGTTGAAGATAAGAGCGGGAGAACTTTGAACGTGGCAACCTGGAACTGGAG 1305
Db 1119 CCAGTTACTTTTGAAGACAAAACGGAAGCCNACTATGACAGAGAAACATGGAGCTGGAG 1178
QY 1306 AAACGAAGCAAGCTCTCTCTGGAACAGAGCGCAAGGAGCAGAGCGGCTTGGCCAGCTG 1365
Db 1179 AAGCGACGCAAGTGTGTGATGGAGCAGCAGAGGGAGGCTGAACGCAAGGCCAGAAA 1238
QY 1366 GAGCGCGGAGCAGCAGAGGAGGAGCTGTAGCGCCAGGAGCAAGAGCGCAAAACAAA 1425

Db 1239 GAGAAGGAAGTGGGAGCGGAAACAGAGAGAACTGCAAGACAGAATGGAGAAGCAG 1298
QY 1426 CTGGAACCTGGGAAGCAACTGGAAGAGCAGCGGGAGCTAGAACGCGCAGAGAGAGGAG 1485
Db 1299 CTGGAGTTGGAGAAACGCTTGGAGAAACAGAGAGAGCTGGAGAGACAGCGGAGGAAG 1358
QY 1486 AGGAGGAAGAAATTTGAGAGCGGAGAGCTGCNAAACGGGAACCTTGAAGGCAACGACAA 1545
Db 1359 AGGAAAGGAGATAGAAAGACGAGAGCGAACAACAGAGCTTTGAGAGACAAACCCCT 1418
QY 1546 CTTGAGTGGGAACGGAATCGAAGGCAAACTTACTAAATCAAAAGAAACAAAGAAACAAG 1605
Db 1419 TTAGAAATGGGAAGAGACTCGGTCCGACAGAGCTGCTCAGTCAGAAGCAGGGAACAAGAA 1478
QY 1606 GACATAGTTGTACTGAAAGCAAAAGAAAGACTTTTGGAAATTTGAATTTAGAAGCTCTAAAT 1665
Db 1479 GACATTTGTCAGGCTGAGCTCCAGAAAGAAAGTCTCCACCTGGAACTGGAGCAGTGAAT 1538
QY 1666 GATAAAAGCATCAACTAGAAGGGAACCTTCAAGATATCAGATGTCGATTGACCAACCAA 1725
Db 1539 GGAACATCATCAGCAGATCTCAGGCGAGACTACAAGATGTCCAATCAGAAAGCAACACAA 1598
QY 1726 AGGCAAGAAATTTAGAGCAGCAAAACAAATCTAGAGAGTTGAGAATTTGCCGAATCAACCAT 1785
Db 1599 AAGACTGAGCTAGAAGTTTGGATTAACAGTGTGACCTGGAAATTTATGNAATCAACAA 1658
QY 1786 CTACAGCAACAAATTAAGGAATCTCAGCAAAATGCTTGGAAAGACTTTATTTCCAGAAAAACAG 1845
Db 1659 CTTCAACAAGAGCTTTAAGGAATATCAAAATAAGCTTATCTATCTGCTCCCTGAGAAGCAG 1718
QY 1846 ATACTCAATGACCAATTAACCAAGTTTCAGCAGAACAGCTTTTGCACAGAGATTCACCTGTT 1905
Db 1719 CTATTAAACGAAAGAAATTTAAACACATGCAGCTCAGTAAACACACCTGATTCAGGGATCAGT 1778
QY 1906 ACACTTAAAGAGCCTTAGAAGCAAAAGAACTAGCTCGGCGACACTACGAGACCAACTG 1965
Db 1779 TTACTTTCAAAAGTCTATCAGAAAGAGAAATTTATGCCAAGACTTTAAGAACAAATTA 1838
QY 1966 GATGAAGTGGAGAAAGAACTAGATCAAAACCTACAGGAGATTTGATTTTCAATTAATCAG 2025
Db 1839 GATGCTCTTGAAGAAAGAACTGCATCTAAGCTCTCAGAAATGGATTCATTTAAACAATCAG 1898
QY 2026 CTGAAGGAACTAAGGAATACACATTAAGCAACAACTCCAGAGCAAAAGTCCATGGAG 2085
Db 1899 CTGAAGAACTCAGAGAAAGCTTATATACACAGCAGTTTAGCCCTTTGAACAACTTCATAAA 1958
QY 2086 GCTGAACGACTGAAACAGAAAGAAACAAAGAACTAGAGATTCATAGAAATTTAGAAAAACAAA 2144
Db 1959 ATCAACCTGACAAATTTGAAGAAATCGAAGAAAGAAATTAGAGCAAAAAAANAANA 2017

RESULT 5

US-09-216-003A-72

; Sequence 72, Application US/09216003A

; Patent No. 6670463

; GENERAL INFORMATION:

; APPLICANT: Mitcham, Jennifer L.

; APPLICANT: Frudakis, Tony N.

; APPLICANT: King, Gordon E.

; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY OF OVARIAN CANCER

; FILE REFERENCE: 210121.462

; CURRENT APPLICATION NUMBER: US/09/216,003A

; CURRENT FILING DATE: 1998-12-17

; NUMBER OF SEQ ID NOS: 310

; SOFTWARE: PatentIn Ver. 2.0

; SEQ ID NO 72

; LENGTH: 2017

; TYPE: DNA

; ORGANISM: Homo sapiens

US-09-216-003A-72

Query Match 9.8%; Score 507.8; DB 3; Length 2017;
Best Local Similarity 55.8%; Pred. No. 7.3e-111;

| Matches 1138; Conservative 0; Mismatches 802; Indels 99; Gaps 5; | |
|--|---|
| QY 175 | GGCGTCGATTAGCAAGTTAAAGTAACAGAAACATGGCTCAGTTTCCAAACACCTTTTGGT 234 |
| Db | |
| QY 9 | GGCTGAGAGCTGCAAGAAAGTCCAGGATCATGATGGCTCAGTTTCCCAACAGCGATGAAT 68 |
| Db | |
| QY 235 | GGCAGCTGGATATCTGGGCCATNACTGTAGAGGAAGAGCGAAGCATGATCAGCAGTTTC 294 |
| Db | |
| QY 69 | GGAGGGCCAAATATTGGGCTATTATCATCTGAAGAACGTAATAGCATGATAAACAAGTTT 128 |
| Db | |
| QY 295 | CATAGTTTAAAGCCAAATATCTGGATTCAATTCTGTGTGATCAAGCTAGAAACTTTTTTTT 354 |
| Db | |
| QY 129 | GATNACTCAACCTTCAGGAGTTACATAACAGGTGATCAAGCCGTACTTTTTTCCTA 188 |
| Db | |
| QY 355 | CAATCTGGTTACTCAACCTGTGTTTATGACAGATATGGGCACTAGCTGACATGAATAT 414 |
| Db | |
| QY 189 | CAGTCAGGTCTGCGGCCCGGTTTATGCTGAAATATGGGCTTATCAGATCTGAACAAG 248 |
| Db | |
| QY 415 | GATGGAAAGTGGATCAAGTGGAGTTTCCATAGCTATGAACCTTATCAAACTGAAGCTA 474 |
| Db | |
| QY 249 | GATGGAAAGTGGACAGCAAGAGTTCTCTATAGCTATGAACCTCATCAAGTTAAAGTTG 308 |
| Db | |
| QY 475 | CAAGGATATCAGTACCTCTGCACTTCCCTCTGTCATGAACAGCAACC---AGTTGCT 531 |
| Db | |
| QY 309 | CAGGGCCACAGCTGCTGTAGTCTCTCTCTCTATCATGAACAACCCCTATGTTCTCT 368 |
| Db | |
| QY 532 | ATTTCTAGGCAACAGATTTGGTATGGAGGTATCGCAGCATGCGCAGCTTACAGCT 591 |
| Db | |
| QY 369 | CCACTAATCTCTGCTGTTTGGGATGGGAAGCATGCCAATCTGTCCATTCATCAGCCA 428 |
| Db | |
| QY 592 | GTGCTCCAGTGGCAATGGGATC-----CAAT 618 |
| Db | |
| QY 429 | TTGCCCTCCAGTGGCACTATAGCAACCCCTGTCTTCTGCTACTTTCAGGGACCATGAT 488 |
| Db | |
| QY 619 | CCAGTTGTTGGAATGTCTCAACCCCTAGTATCTTCTGTTCCCAACAGCAGCTGTCGCCCC 678 |
| Db | |
| QY 489 | CTTCCCTTAATGATGCTGTCTCCCTAGTCTCTGTTAGTACATCTCTTACCAAT 548 |
| Db | |
| QY 679 | CTGGCTAAAGGGGCTCCCTGTTATACAACTCTGCTGCTGATTTGCTCATCTCGAGCC 738 |
| Db | |
| QY 549 | GGAACTGCCAGTCTCATTCAGCCCTTATCCATTCCTTATCTTCTTCAACATTTGCTCAT 608 |
| Db | |
| QY 739 | ACATTGCCAAGAGTCTTCTCTTATGATCTGTTCCAGGGTCACAACTTAAACACTAA 798 |
| Db | |
| QY 609 | GCATCATCTTACAGCCCTGATGATGGGAGGATTTGGTGGTCTGATATCCAGAGGCCAG 668 |
| Db | |
| QY 799 | TTACAAAGGCACAGTCAATTTGATGTGGCCAGTGTCCACACAG-----841 |
| Db | |
| QY 669 | TCTCTGATTTAGGATCTAGTACTCACTTCTCTCACTGCTTCCCTCTCAGGGAAC 728 |
| Db | |
| QY 842 | -----TGGCAGAGTGGGCTGTTCTCAGTCATCAAGCTGAATATACAGG 885 |
| Db | |
| QY 729 | TCACCTAAGACAGGGACCTCAGAGTGGGAGTTCTCAGGCTTCAAGATTTAAAGTATCGG 788 |
| Db | |
| QY 886 | CAATTATCAATAGTATGACAAACATCATGATGGACACTTAAACAGTCCCAAGCAAGA 945 |
| Db | |
| QY 789 | CAAAAATTTAATAGTTAGACAAAGGATGAGCGGATACCTCTCAGGTTTCAAGCTAGA 848 |
| Db | |
| QY 946 | ACTATTCTTATGAGTCAAGTTTACCACAGGCTCAGCTGGCTTCAATATGGAATCTTTCT 1005 |
| Db | |
| QY 849 | AATGCCCTTCTTCTGATCAATCTCTCTCAACTCAGCTAGTACTATTTGGACTCTGGCT 908 |
| Db | |
| QY 1006 | GACATTCATCAAGATGGAACCTTACAGCAGAGGAATTTATCTCTGGCAATGCACTCAT 1065 |
| Db | |
| QY 909 | GACATCGATGGTACCGACAGTTGAAAGCTGAAGAAATTTATTTCTGGCGATGCACTCACT 968 |
| Db | |
| QY 1066 | GATGTACTATGTCTGCCACCACTGCCACCTGCTCTGCTCCAGAAATACATTTCCACCT 1125 |
| Db | |
| QY 969 | GACATGCCAAAGCTGACACGCCATACCACTGAGTTGCTCTCCGAGCTTGTCCCTCA 1028 |
| Db | |
| QY 1126 | TCITTTAGAGAGTTCGATCTGGCAGTGGTATATCTGTCTAAGCTCAACATCTGTAGAT 1185 |
| Db | |
| QY 1029 | TCITTCAGAGGGGAAGCAAGTTGAT-----TCTGTAAAGAACTCTGCTTCATAT 1082 |
| Db | |

RESULT 6

US-09-667-857-72
; Sequence 72, Application US/09667857
; Patent No. 6699664
; GENERAL INFORMATION:
; APPLICANT: Mitcham, Jennifer L.
; APPLICANT: King, Gordon E.

| | |
|---------|---|
| QY 1186 | CAGAGGCTACAGAGGAAACCACTTTTAGAAGATGAACAACAACAAATTAGAAAAGAAATTA 1245 |
| Db | |
| QY 1083 | CAGAAAAACAAGAGAGAGCCT-----CAGAAGAACTG 1118 |
| Db | |
| QY 1246 | CCTGTAAAGCTTTGAAGATAAGAGCGGGAGAACTTTGAACGTGGGCACCTCGAACCTGGAG 1305 |
| Db | |
| QY 1119 | CCAGTTACTTTTGGAGCAAAACGGAAGCCAACTATGAACGAGGAACAATGGAGCTGGAG 1178 |
| Db | |
| QY 1306 | AAACGAGGCAAGCTCTCTGGAACAGCAGCGCCAAAGGAGCAGAGCGCTGGCCCAAGCTG 1365 |
| Db | |
| QY 1179 | AAGCCAGCGCAAGTGTGTATGAGCAGCAGAGAGGAGGCTGAACGCCAAGCCCAAGAA 1238 |
| Db | |
| QY 1366 | GAGCGGCGGAGCAGAGAGAGGAGAGCGTGAAGCGCCAGGAGCAGAGCGCCAAAAGCAAA 1425 |
| Db | |
| QY 1239 | GAGAGGAGAGTGGGAGCGGAAACAGAGAGAACTGCAAGAGCAAGAAATGGAAGAGCAG 1298 |
| Db | |
| QY 1426 | CTGGAACTGGAGAACTCTGGAAGAGCAGCGGAGCTAGAACGGCAGAGAGAGGAGGAG 1485 |
| Db | |
| QY 1299 | CTGGAGTTTGGAGAAACGCTTGGAGAAACAGAGAGAGCTGGAGAGACAGCGGGAGAGAG 1358 |
| Db | |
| QY 1486 | AGGAGAAAGAAATTCAGAGGCGAGAGGCTGCAAAAACGGGAACTTCAAAAGGCAACACAA 1545 |
| Db | |
| QY 1359 | AGGAGAAAGAGATAGAAAGACGAGAGGAGCAAAAACAGAGCTTGAAGACAAACGCCGT 1418 |
| Db | |
| QY 1546 | CTTGTAGTGGGAAACGGAATCGAAAGGCAAGAACTACTAAATCAAGAAACAAAGAACAGAG 1605 |
| Db | |
| QY 1419 | TTAGATGGGAAGAGCTCCGTCGGCAGGAGCTGCTCAGTCAGAAGACCAGGGAACAAGAA 1478 |
| Db | |
| QY 1606 | GACATAGTTGTACTGAAACAAAGAAAGAGACTTTTGGAAATTTGAAATTTAGAGCTTAAT 1665 |
| Db | |
| QY 1479 | GACATTTGTGAGCTGAGCTCCAGAAAGAAAGTCTCCACTTGAAGTGAAGCAGTGAAT 1538 |
| Db | |
| QY 1666 | GATAAAGCATCACTAGAGGGAACCTTCAAGATATCAGATGTGATTTGACCAACCCAA 1725 |
| Db | |
| QY 1539 | GGAAAAACATCAGCAGATCTCAGGACAGCTACAGATGTCCAATTCAGAAAGCAACACAA 1598 |
| Db | |
| QY 1726 | AGCAGAAATTTAGAGACCAACAAATCTAGAGAGTTTGAGAAATTTGCCGAATTTAGCAAT 1785 |
| Db | |
| QY 1599 | AGACTGAGCTAGAGTTTGGTAAACAGTGTGACCTGGAATTTATGGAATTCAAACAA 1658 |
| Db | |
| QY 1786 | CTACAGCAACAAATTACAGGAATCTCAGAAATCTTGGAAAGTGTATTTTCAGAAAGAACAG 1845 |
| Db | |
| QY 1659 | CTTCAACAGAGCTTAAGGAATATCAAAATTAAGCTTATCTATCTGGTCCCTGGAAGCAG 1718 |
| Db | |
| QY 1846 | ATACTCAATGACCAATTAACAAAGTTTCAGCAGAAACAGTTTGACAGAGATTCACCTTGT 1905 |
| Db | |
| QY 1719 | CTATTAAACGGAAGAAATTAACAAACATGACAGCTCAGTAAACACACTGATTCAGGATCAGT 1778 |
| Db | |
| QY 1906 | ACACTTAAAGAGCCTTTAGAAACAAAGAACTAGCTCGGAGCAACCTTACGAGACCAACTG 1965 |
| Db | |
| QY 1779 | TTACTTCTAATAAAGTCTATCAGAAAGGAAGAAATTTATGCCAAGACTTAAAGAAACAATTA 1838 |
| Db | |
| QY 1966 | GATGAAGTGGAGAAAGAACTAGATCAAAACCTACAGAGATTTGATATTTTCAATATACAG 2025 |
| Db | |
| QY 1839 | GATGCTCTTGAAGAAAGAACTGATCTAAGCTCTCAGAAATTTGATTTCAATTAACATCAG 1898 |
| Db | |
| QY 2026 | CTGAAGAACTAGAGAAATACAAATNAAGCAACAACTCCAGAAAGCAAAAGTCCATGGAG 2085 |
| Db | |
| QY 1899 | CTGAAGAACTCAGAGAAAGCTATAATACACAGCTTAGCCCTTGAACAACTTCATATAA 1958 |
| Db | |
| QY 2086 | GCTGAAGCAGCTGAAACAGAAAGAAACAAAGAACGAAAGATCATAAGATTTAGAGCAAAA 2144 |
| Db | |
| QY 1959 | ATCAACGCTGACAAATTTGAAGGAATTCGAAGAAAGAAAGATTTAGAGCAAAAAGAAAAA 2017 |
| Db | |

```
/ APPLICANT: Algate, Paul A.
/ APPLICANT: Fling, Steven P.
/ APPLICANT: Retter, Marc W.
/ APPLICANT: Fanger, Gary Richard
/ APPLICANT: Reed, Steven G.
/ APPLICANT: Vedwick, Thomas S.
/ APPLICANT: Carter, Darrick
/ TITLE OF INVENTION: POSITIONS AND METHODS FOR THE THERAPY AND
/ TITLE OF INVENTION: DIAGNOSIS OF OVARIAN CANCER
/ FILE REFERENCE: 210121.462C5
/ CURRENT APPLICATION NUMBER: US/09/667,857
/ CURRENT FILING DATE: 2000-09-20
/ NUMBER OF SEQ ID NOS: 455
/ SOFTWARE: FastSeq for Windows Version 3.0
/ SEQ ID NO 72
/ LENGTH: 2017
/ TYPE: DNA
/ ORGANISM: Homo sapien
US-09-667-857-72

Query Match          9.8%; Score 507.8; DB 3; Length 2017;
Best Local Similarity 55.8%; Pred. No. 7.3e-111;
Matches 1138; Conservative 0; Mismatches 802; Indels 99; Gaps 5;

QY 175 GCGCTGATTAGCAAGGTAAAGTAAACAGAACCATGGCTCAGTTTCCAAACACCTTTTGGT 234
DB 9 GCGCTGAGAGCTGCAAGAAGAAGTCCAGGATCATGATGGCTCAGTTTCCCAAGCGATGAAT 68

QY 235 GCGAGCTGGATATCGGGCCCACTAGTGTAGGGAAGAGCGAAGCATGATCAGCAGTTC 294
DB 69 GGAGGGCCAAATATGFGGCTATTACATCTGAAGAAGCTAAGCATGATGAACAGTTT 128

QY 295 CATAGTTTAAAGCCAAATATCTGGATTCAATTAAGTGTGATCAAGCTAGAACTTTTTTTT 354
DB 129 GATTAACCTCAACCTTCAGGAGGTATACATAACAGGTGATCAAGCCGTACTTTTTCCTA 188

QY 355 CAATCTGGGTACTTCAACCTGTTTGTAGCACAGATATGGGCTAGCTGACATGAATAAT 414
DB 189 CAGTCAGGTCTGCGGCCCGGTTTTAGCTGAAATATGGGCTTATCAGATCTGAACAAG 248

QY 415 GATGGAAGATGGATCAAGTGAGGATTTCCATAGCTATGAACCTTATCAACTGAAGCTA 474
DB 249 GATGGAAGATGGACCAAGAGTTCTCTATAGCTATGAACCTATCAAGTTAAAGTTG 308

QY 475 CAAGGATATCAGTACCTCTGCACTTCCCTCTGTCATGAACACAGCAAC---AGTTGCT 531
DB 309 CAGGGCCACAGCTGCTGTAGTCCCTCCTCTATCATGAACAAACCCCTATGTTCTCT 368

QY 532 ATTTCTAGCGCACCGATTTGGTATGGGAGGTATGCCAGCATGCCACGGCTTACAGCT 591
DB 369 CCACTAATCTCTGCTCGTTTGGGATGGGAAGCATGCCCAATCTGTCCATTATCAGCCA 428

QY 592 GTTGCTCCAGTCCCATGGATC-----CATTT 618
DB 429 TTGCTCCAGTTGCACCTATAGCAACCCCTGTGCTTCTGCTACTTTCAGGACCATGATTT 488

QY 619 CCAGTTGTTGGAATGTCTCCAAACCTAGTATCTTCTGTTCCCAAGCAGCTGTGCCCC 678
DB 489 CCTCCCTAATGATGCTGCTCCCTTAGTGCTTCTGTTAGTACATCTCATTTACCAAT 548

QY 679 CTGGCTTAAGGGGCTCCCTGTTTATACAACTTCTGCTGCTGCTATCTGCTATCTGAGCC 738
DB 549 GGAACCTGCCAGTCTCATTCAGCCCTTTATCCATTCTCTTATTTCTTCTCAACATTTGCC 608

QY 739 ACATTGCCAAAGAGTTCTTCTTTAGTATGTTCTGCTCCAGGGTCCAACTTAAACACTAA 798
DB 609 GCATCATCTTACAGCTGATGATGGAGGATTTGGTGGTGTGCTAGTATCCAGAGGCCAG 668

QY 799 TTACAAAGGCACAGTCATTTGATGTGGCCAGTGTCCCAACCG-----841
DB 669 TCTCTGATTGTTAGGATCTAGTACTCAACTTCTCAACTGCTTCCCTCTCAGGGAAC 728

QY 842 -----TGGCAGAGTGGGCTGTTCTCTCAGTCATCAAGACTGAATATACAGG 885
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DB 729 TCACCTAAGACAGGAGCCTCAGAGTGGCAGTTCCTCAGCCTTCAAGATTAAAGTATCGG 788
QY 886 CAAATTATTCAATAGTATGACAAAACCTATGAGTGGACACTTAAACAGGTCCTCCCAAGCA 945
DB 789 CAAAAATTTAATAGTCTAGACAAAGGCATGAGCGATACCTCTCAGGTTTTCAAGCTAGA 848
QY 946 ACTATCTTATGCAAGTCAAGTTTACCAGAGGCTCAGTGGCTTCAATATGGAATCTTCT 1005
DB 849 AATGCCCTTCTTCAAGTCAAACTCTCTCAAACTCAGTAGCTACTATTGGAATCTGGCT 908
QY 1006 GACATTGATCAAGATGGAAACTTACACGAGAGGAATTTATCTCGGCAATGCACTCATTT 1065
DB 909 GACATCGATGGTGACGACAGTTGAAGCTGAAGAAATTTATTTCTGGCCATGACCTCACT 968
QY 1066 GATGTAGCTATGTCTGGCCAAACCACTGCCACCTGTCTGCTCTCCAGAAATACATTCAC 1125
DB 969 GACATGGCCAAAGCTGGACAGCACTACCACTGACGTTTGGCTTCCGAGCTTGTCTCCCA 1028
QY 1126 TCTTTTGAAGAGTTCGATCTGGCAGTGGTATATCTGTCTATAGCTCAACATCTGTAGAT 1185
DB 1029 TCTTTTCAAGAGGGGAAAGCAAGTTTGTAT-----TCTGTTAATGGAACCTTGCCTT 1082
QY 1186 CAGAGGCTTACAGAGAGAACCAAGTTTGTAGAGATGAACAACAATTTAGAAAAGAAATTA 1245
DB 1083 CAGAAAAACAAAGAAAGAGAGCT-----CAGAAAGAACTG 1118
QY 1246 CCGTGTAAACCTTTGAAGATAAGAAAGCGGAGAACTTTTGAACGTGGCAACCTTGAAC 1305
DB 1119 CAGATTACTTTTGTAGGACAAACCGAAAGCCAACTATGAACGAGGAAACATGAGCTGG 1178
QY 1306 AAACGAAAGCAAGCTCTCTGTGAAACAGCAGCGCAAGGAGAGGAGCGCTGCCCCAGCTG 1365
DB 1179 AAGCGACGCCAAGTTTGTGTGAGCAGCAGAGAGGAGGCTGAAACGCAAAAGCCAGAAA 1238
QY 1366 GAGCGGCGGAGCAGCAGAGAGGAGGAGCGTGTAGCGCCAGGAGCAAGAGCGCAAAAGACA 1425
DB 1239 GAAAGGAGAGTGGGAGCGGAAACAGAGAGAACTGTCAAGACAAAGATGGAAGAGCAG 1298
QY 1426 CTGGAACCTGGGAAGCAACTCGAAAGCAGCGGAGCTAGAAACGCGCAGAGAGAGGAGG 1485
DB 1299 CTGGAGTTGGAGAAACGCTTGGAGAAACAGAGAGAGCTGGAGAGACAGCGGAGGAAG 1358
QY 1486 AGGAGAAAGAAATTTGAGAGGCGAGAGCTGCAAAACGGGAACCTTGAAGGCAACGACAA 1545
DB 1359 AGGAGAAAGGAGATGAAAGACGAGAGCGCAAAACAGGAGCTTTGAGAGACAAAGCCGT 1418
QY 1546 CTTGAGTGGGAACGGAATCGAAGGCAAACTACTAAATCAAAGAAACAAAGAAACAAGAG 1605
DB 1419 TTAGAAATGGGAAGAACTCCGTCGGCAGGAGCTGCTCAGTCAGAACCCAGGGAACAGNA 1478
QY 1606 GACATAGTTGTACTGAAAGCAAAAGAAAGACTTTTGGAAATTTGAATTTAGAGCTCTAAT 1665
DB 1479 GACATTGTGAGCTGAGCTCCAGAAAGAAAGTCTCCACCTGGAACTGGAGCAGTGAAT 1538
QY 1666 GATAAAAGCATCAACTAGAGGGAACCTTCAAGATATCAGATGTGATGTGATGACCCCAA 1725
DB 1539 GGAAACATCATCAGCAGATCTCAGGCAAGCTACCAAGATGTCCAATCAGAAAGCAACACAA 1598
QY 1726 AGGCAAGAAATTTGAGAGCAAAACAAATCTAGAGAGTTGAGAAATTTGCCAAATCAACCA 1785
DB 1599 AAGACTGAGCTAGAAAGTTTGGATTAACAGTGTGACCTGGAAATTTATGGAATCAACAA 1658
QY 1786 CTACAGCAACAAATTAAGGAATCTCAGCAAAATGCTTGGAAAGACTTTATTCAGAAACAG 1845
DB 1659 CTTCAACAAGAGCTTAAAGGAATATCAAAATAAGCTTATCTATCTGCTCCCTGAGAAGCAG 1718
QY 1846 ATACTCAATGACCAATTTAAACAAAGTTTCAAGAGACAGTTTGCACAGAGATTCATTTGT 1905
DB 1719 CTTATTAACGAAAGAAATTTAAACAAATGTCAGCTCAGTAACACACCTGATTCAGGGAT 1778
QY 1906 ACACCTTAAAGAGCCTTAGAAGCAAAAGAACTAGCTCGGACGACCTACGAGACCAACTG 1965
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Db 1779 TTACTTCATAAAAGTCATCAGAAAAGGAGAAATATATGTCCTTAAAGACATTTAAAGCAATTA 1838
Qy 1966 GATGAAGTGGAGAAAGAACTAGATCAAACTACAGGAGATGATATTTTCAATATACAG 2025
Db 1839 GATGCTCTTGAAGAAAGAACTGATCTAAGCTCTCAGAAATGATTTCAITTAACAATCAG 1898
Qy 2026 CTGAAGGAATAGAGAAATACAAATAGCAATAGCAACACTCCAGAAAGCAAAAGTCCATGGAG 2085
Db 1899 CTGAAGGAATCTCAGAAAGCTATATACACAGAGTTAGCCCTTGAACAACTTCATATAA 1958
Qy 2086 GCTGAACGACTGAAACAGAAAGAAACAAAGAACGAAAGATCATAGAAATAGAAAAACAAA 2144
Db 1959 ATCAAAGCTGACAAATTTGAAGAAATCGAAGAAAGAAAGATTAGAGCAAAAAA 2017

RESULT 7

US-10-198-053-72

; Sequence 72, Application US/10198053

; Patent No. 6858710

; GENERAL INFORMATION:

; APPLICANT: Bangur, Chaitanya S.

; APPLICANT: Retter, Marc W.

; APPLICANT: Fanger, Gary R.

; APPLICANT: Hill, Paul

; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY

; FILE REFERENCE: 210121.462C9

; CURRENT APPLICATION NUMBER: US/10/198,053

; CURRENT FILING DATE: 2002-07-17

; NUMBER OF SEQ ID NOS: 624

; SOFTWARE: FastSeq for Windows Version 4.0

; SEQ ID NO 72

; LENGTH: 2017

; TYPE: DNA

; ORGANISM: Homo sapiens

US-10-198-053-72

Query Match 9.8%; Score 507.8; DB 3; Length 2017;

Best Local Similarity 55.8%; Pred. No. 7.3e-111;

Matches 1138; Conservative 0; Mismatches 802; Indels 99; Gaps 5;

Qy 175 GCGCTCGATTAGCAAGCTAAAGTAAACAGAACCATGGCTCAGTTTCCACACCTTTTGGT 234
Db 9 GCGTGAGAGCTGCAAGAAAGAGTCAGGATCATGATGGCTCAGTTTCCACAGCGATGAAT 68
Qy 235 GCGAGCCTGGATATCTGGGCCATAAATCTGTAGAGGAAAGAGCGAAGCATGATCAGCAGTTC 294
Db 69 GGAGGGCCAATATGTGGGCTATTACATCTGAAGAACGTACTAAGCATGATAAACAGTTT 128
Qy 295 CATAGTTTAAAGCCAATATCTGGATTCATTTAGTGGTATCAGCTAGAACTTTTTTTT 354
Db 129 GATAACCTCAAACTTCAGGAGGTTACATAACAGGTGATCAAGCCCGTACTTTTTTCCCTA 188
Qy 355 CATCTGGGTACCTCAACCTGTTTATAGCAGAGATATGGGCATCTAGCATGAATTAAT 414
Db 189 CAGTCAGTCTGCGGCGCCGGTTTTAGCTGAAATATGGGCTTTATCAGATCTGAAACAG 248
Qy 415 GATGGAAGATGGAATCAAGTGGAGTTTTCATAGCTATGAACCTTATCAAACTGAAGCTA 474
Db 249 GATGGGAAGATGGACCAAGAGAGTCTCTATAGCTATGAACCTTATCAAGTTAAAGTTG 308
Qy 475 CAAGGATATCAGCTACCTCTGCACTTCCCTCTGCACTTCCCTCTGCACTTGAAGAACAGCA 531
Db 309 CAGGGCCAAACAGCTGCTGTAGTCTCCCTCTATCATGAAGAACACCCCTTATGTTCTCT 368
Qy 532 ATTTCTAGCGCACCGAGATTTGGTATGGAGGTATCGCCAGCATGCGCACCGCTTACAGCT 591
Db 369 CCAGTAAATCTCTGCTCGTTTTGGGATGGAAAGCATGCCCATCTGTCCATTCATCAGCCA 428
Qy 592 GTTCTCAGTGCCCAATGGGATC-----CATT 618
Db 429 TTGCTCCAGTTGACCTATAGCAACACCCCTTGTCTTCTGCTACTTTCAGGGACCGATATT 488


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Db      1539  GGAAGAAATCATCAGCAGATCTCAGCAGACTACAGATGTCCAATCAGAAAGCAACACAA 1598
Qy      1726  AGCGAAGAAATTTAGAGAGCAACAACAAATCTAGAGAGTTGAGAGATTCGCCAAATCAACCAAT 1785
Db      1599  AAGACTGAGCTAGAAGTTTGGATTAACAAGTGTGACCTGGAAATTTATGGAAATCAACAA 1658
Qy      1786  CTACAGCAACAATTTACAGGAATCTCAGCAAAATCTCTTGGAAAGACTTATTTCCAGAAAAACAG 1845
Db      1659  CTTCAACAAGAGCTTAAGGAATATCAAAATAGCTTTATCTATCTGTGCTCCCTGAGAAGCAG 1718
Qy      1846  ATACTCAATGACCAATTAACAAAGTTTCAGAGAACAGATTTGCACAGAGATTCACATTGTT 1905
Db      1719  CTTATTAACGAAAGATTTAAATAACATGACGCTCAGTAAACACACCTGATTCAGGGATCAGT 1778
Qy      1906  ACACCTTAAAGAGCCCTTGAAGCAAAAGAACTAGCTCGGCAGCAGCACCTACGAGACCAACTG 1965
Db      1779  TTACTTCATAAAAGTCATCAGAAAGGAAGAAATTTATGCGCAAGACTTAAAGAACAAATTA 1838
Qy      1966  GATGAAGTGGAGAAAGAACTAGATCAAAACTACAGGAGATTTGATATTTTCAATATATCAG 2025
Db      1839  GATGCTCTTTGAAAAGAAAGAACTGCATCTAAGCTCTCAGAAATGGAATTCATTTAAACAATCAG 1898
Qy      2026  CTGAAGGAATTAAGGAATATACAAATAGCAACAACTCCAGAACGAAAGTCCATGGAG 2085
Db      1899  CTGAAGGAATCTCAGAGAAAGCTATATACACAGCAGTTAGCCCTTTGAAACAACCTTCATAAA 1958
Qy      2086  GCTGAACGACTCAAAACAGAAAGAAACAAAGAACGAAAGACTATAGAAATTTAGAAAAACAAA 2144
Db      1959  ATCAAAAGTGCACAAATTTGAAGGAATCGAAAGAAAGAAATTTAGAGCAAAAAAANAANA 2017
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RESULT 8

US-09-827-271-72

; Sequence 72, Application US/09827271

; Patent No. 6962980

; GENERAL INFORMATION:

; APPLICANT: Retter, Marc W.

; APPLICANT: Fanger, Gary R.

; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND

; TITLE OF INVENTION: DIAGNOSIS OF OVARIAN CANCER

; FILE REFERENCE: 210121.462C6

; CURRENT APPLICATION NUMBER: US/09/827,271

; CURRENT FILING DATE: 2001-04-04

; NUMBER OF SEQ ID NOS: 461

; SOFTWARE: FastSeq for Windows Version 3.0

; SEQ ID NO 72

; LENGTH: 2017

; TYPE: DNA

; ORGANISM: Homo sapien

US-09-827-271-72

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Query Match          9.8%; Score 507.8; DB 3; Length 2017;
Best Local Similarity 55.8%; Pred. No. 7.3e-111;
Matches 1138; Conservative 0; Mismatches 802; Indels 99; Gaps 5;

Qy      175  GCGTGCATTTAGCAAGTAAAAGTAAACAGAACCATGGCTCAGTTTCCACACCTTTTGGT 234
Db      9    GCGTGAGAGCTGCAAGAAAGAGTCAGGATCATGTGGCTCAGTTTCCACAGCGATGAAT 68
Qy      235  GCGAGCTGGATATCTGGGCCATACTGTAGAGGAAGAGCGNAGCATGATCAGCAGTTTC 294
Db      69  GGAGGGCCAAATATGTGGGCTATTACATCTGAAGAACGTAATAAGCATGATAACAGTTT 128
Qy      295  CATAGTTTAAAGCCAAATATCTGGATTCAATTACTGTGTGATCAAGCTAGAAACTTTTTTTT 354
Db      129  GATAACCTCAAACCTTCAGAGGTTTACATAACAGGTGATCAAGCCCGTACTTTTTTCCTA 188
Qy      355  CAATCTGGGTTACCTCAACCTGTTTTTACACAGATATGGGCACTAGCTGACATGAATAAT 414
Db      189  CAGTCAGGTCTGCGGCCCGGTTTTTAGCTGAAATATGGGCTTTATCAGATCTGAACAAG 248
Qy      415  GATGNAGAAATGGATCAAGTGGAGTTTTTCCATAGCTATGAAACTTATCAAACTGAAGCTA 474
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Db      249  GATGGGAAGATGGACCAAGAGAGTTCTATAGCTATGAAACTCATCAAGTTAAAGTTG 308
Qy      475  CAAGGATATCAGCTACCCCTCTGCACCTCCCTCTGTCATGAAACAGCAACCC---AGTTGCT 531
Db      309  CAGGCGCAACAGCTGCTGTAGTCTCTCTCTATCATGAACAACCCCTATGTTCTCT 368
Qy      532  ATTTCTAGCGCACCAAGCATTTGGTATGGAGGATATCGCAGCATGCCACCGCTTACAGCT 591
Db      369  CCACATAATCTCTGCTCGTTTTGGGATGGGAAGCATGCCCAATCTGTCCATTTCATCAGCCA 428
Qy      592  GTTGCTCCAGTGCCCAATGGATC-----CAATT 618
Db      429  TTGCTCCAGTTGCACTTATAGCAACCCCTTGTCTTCTGTACTTTCAGGGAGCCAGTATT 488
Qy      619  CCAGTTGTTGGAATGCTCCCAACCTTAGTATCTCTGTTTCCACAGCAGCTGTGCCCC 678
Db      489  CTTCCCTTAATGATGCTGCTCCCTCTAGTGCCTTCTGTTAGTACATCCTCATTACCAAT 548
Qy      679  CTGGCTAACGGGGCTCCCTCTGTTATACAACTCTGCTGCAATTTGCTCATCCTTGAGCC 738
Db      549  GGAACCTGCAGTCTCATTCAGCCCTTATCCCATTTCTTATCTTCTTCAACATTGGCTCAT 608
Qy      739  ACATTTGCCCAAGAGTTCTTCTTTAGTAGATCTGGTCCAGGGTCAACAATAACAACATAA 798
Db      609  GCATCATCTTACAGCTGATGATGGAGGATTTGGTGGTGTAGTATCCAGAAAGGCCAG 668
Qy      799  TTACAAAAGGCACAGCTCATTTGATGTGCCAGTGTCCACCCAG----- 841
Db      669  TCTCTGATTGATTAGGATCTAGTAGCTCAACTTCTCAACTGCTTCCCTCTCAGGGAAC 728
Qy      842  -----TGGCAGAGTGGGCTGTTCTCTAGTCAATCAAGACTGAAATACAGG 885
Db      729  TCACCTTAAGACAGGGACCTCAGAGTGGCGAGTTCTCAGCCCTTCAAGATTAAAGTATCGG 788
Qy      886  CAATTTATTCATAGTCATGACAAAATATAGTGGACACTTAAACAGGTCCCAAGCAAGA 945
Db      789  CAAAAATTTAATAGTCTAGACAAAAGCATGAGCGGATACCTCTCAGGTTTTTCAGCTAGA 848
Qy      946  ACTATTTCTATGCACTCAAGTTTACACAGGCTCAGCTGGCTTCAATATGGAATCTTCT 1005
Db      849  AATGCCCTTCTCAGTCAAAATCTCTCAAACTCAGTAGCTACTATTTGGACTCTGGCT 908
Qy      1006  GACATTTGATCAAGATGGAAAACTTACAGAGAGGAATTTATCCTGGCAATGCACTCAT 1065
Db      909  GACATCGATGGTGCAGCAGTTGAAAGCTGAAAGAAATTTATTTCTGGCGATGCACTCACT 968
Qy      1066  GATGTAGTATGTCTGGCCCAACCACTGCCACCTGCTCTGCTCCAGAAATACATTCACCT 1125
Db      969  GACATGGCCAAAGCTGGACAGCCACTACCATGAGCTTGGCTCCCGAGCTTGTCCCTCCA 1028
Qy      1126  TCTTTTGAAGAGTTTCGATCTGGCAGTGGTATATCTGTATAGCTCAACATCTGTAGAT 1185
Db      1029  TCTTTTCAAGGGGGAAGCAAGTTGAT-----TCTGTTAATGGAACCTCGCTTTCATAT 1082
Qy      1186  CAGAGGCTACCGAGGAACCGAGTTTGAAGATGAACAACAATTAGAAAAAGAAATTA 1245
Db      1083  CAGAAAAACACAGAAAGAGAGCCT-----CAGAAAGAAACCTG 1118
Qy      1246  CCGTTAAACGTTTGAAGATAAGAAAGCGGAGAACTTTTGAACGTGGCAACCTCGAACTGGAG 1305
Db      1119  CAGTTTACTTTTGAAGACAAACGGAAAGCCAACTATGAACGAGGAAACATGAGAGCTGGAG 1178
Qy      1306  AAACGAAGGCAAGCTCTCTCGAAACAGCAGCGCAAGGAGGAGGCGCTGCGCCAGCTG 1365
Db      1179  AAGCGACGCCAAGTGTGTATGGAGCAGCAGAGAGGGAGGCTGAAACGCAAGCCAGAAA 1238
Qy      1366  GAGCGGGCGGAGCAGAGGAGGAGCGTGAAGCCAGGAGCAGAGCGCAAGGCAAGAAACAA 1425
Db      1239  GAGAAAGGAAGTGGGAGCGGAAACAGAGAGAACTTGAAGAGCAAGAAATGGAAGAGCAG 1298
Qy      1426  CTGGAACCTGGAGAAAGCAACTGGAAGAGCAGGGGAGCTAGAACGCGCAGAGAGAGGAG 1485
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| 1299 | CTGGAGTTGGAGAAACGCTTTGGAGAAACAGAGAGAGCTGGAGAGACNGCGGAGGAGAG | 1358 |
| 1486 | AGGAGGAAAGAAATTTAGAGCGGAGAGGCTGCAAAACGGGAACTTTGAAAGGCAACGACAA | 1545 |
| 1359 | AGGAGAAAGGAGATAGAAAGACGAGAGGCAGCAAAACAGGAGCTTGAGAGACAAACGCGCT | 1418 |
| 1546 | CTTGAGTGGGAAACGGAACTCGAAGCGCAGAGACTACTAAATCAAAAGAAACAAGAACAAAGAG | 1605 |
| 1419 | TTAGAAATGGGAAAGACTCCGCTCGGCGAGGAGCTGCTCAGTCAGAGACCCAGGGAACAAGAA | 1478 |
| 1606 | GACATAGTTGTACTGAAAGCAAAAGAAAAGACTTTTGGAAATTTGAAATTTAGAAGCTCTAAAT | 1665 |
| 1479 | GACATTTGTCAGGCTGAGCTCCAGAAAGAAAGTCTCCACCTGGAACTGGAAAGCAGTGAAT | 1538 |
| 1666 | GATAAAAGGATCAACTAGAAAGGGAACCTTCAAGATATCAGATGTGCAATTTGACCAACCCAA | 1725 |
| 1539 | GGAAAACATCAGCAGATCTCAGGCGAGACTACAAGATGTCCAAATTCAGAAAAGCAAAACAA | 1598 |
| 1726 | AGGCAAGAAATTCGAGAGCACAAAACAAATCTAGAGAGTTGAGAAATTCGCCGAAATCACCCAT | 1785 |
| 1599 | AAGACTGAGCTAGAGTTTGGATAAACAGTGTGACCTGGAAATTTAGGAAATCAACAA | 1658 |
| 1786 | CTACAGCAACAATTTACAGGAATCTCAGCAAAATGCTTTGGAAGACTTATTTCCAGAAAAACAG | 1845 |
| 1659 | CTTCAACAAGAGCTTAAGGAATATCAAAATAAGCTTATCTATCTGTGCTCCCTGGAAGACAG | 1718 |
| 1846 | ATACTCAATGACCAATTTAAACAAGTTTCAGCAGAACAGTGTTCACACAGAGATTTCACTTGTT | 1905 |
| 1719 | CTATTAAACGAAAGAAATTTAAAAAACATCGAGCTCAGTAAACACACTGATTCAGGGATCAGT | 1778 |
| 1906 | ACACTTAAAAGAGCCTTTAGAAGCAAAAAGAACTAGCTCGGCAGCACCTACGAGACCAACTG | 1965 |
| 1779 | TTACTTCAATAAAGTCAACAGAAAGGAAAGATTATGCCAAAGACTTAAAGAACAAATTA | 1838 |
| 1966 | GATGAAGTGGAGAAAGAAACTAGATCAAAACTACAGGAGATTTGATATTTTCAATAATTCAG | 2025 |
| 1839 | GATGCTCTTGAAAAGAAACATGCAATCTTAAGCTCTTCGAGAAATGGGAATCATTTAAACAATTCAG | 1898 |
| 2026 | CTGAAGCAACTTAAGAGAAATACACAAATAAGCAACAACTCCAGAGCAAAAAGTCCCATGGAG | 2085 |
| 1999 | CTGANGGAATCAGAGAAAGGCTATTAATACACAGCAGTTAGCCCTTGAACAACTTCATAAA | 1958 |
| 2086 | GCTGAACGACTGAAACAGAAAGAACAGAACGAAAGATCATAGAAATTTAGAAAAACAAAA | 2144 |
| 1959 | ATCAAGCTGCAAAATTTGAAGGAATCGAAGAAAAAGATTTAGAGCAAAAAAAGAAAAA | 2017 |

RESULT 9

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US-09-513-999C-27927
; Sequence 27927, Application US/09513999C
; Patent No. 6783961
; GENERAL INFORMATION:
; APPLICANT: Dumas Milne Edwards, J.B.
; APPLICANT: Duclert, A.
; APPLICANT: Giordano, J.Y.
; TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
; Patent No. 6783961
; FILE REFERENCE: 59.US2.REG
; CURRENT APPLICATION NUMBER: US/09/513,999C
; CURRENT FILING DATE: 2000-02-24
; PRIOR APPLICATION NUMBER: US 60/122,487
; PRIOR FILING DATE: 1999-02-26
; NUMBER OF SEQ ID NOS: 36681
; SOFTWARE: Patent.pm
; SEQ ID NO 27927
; LENGTH: 174
; TYPE: DNA
; ORGANISM: Homo sapiens
; US-09-513-999C-27927

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Query Match      3.3%; Score 174; DB 3; Length 174;
Best Local Similarity 100.0%; Pred. No. 1.2e-31;
Matches 174; Conservative 0; Mismatches 0; Indels
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| Qy | 3936 | TTCCCCCACCCTTTGCACAGGTGCTTTCAATAGCTTTTAAATATATTTTTTAAATATATATTT | 3995 |
| Db | 1 | TTCCCCCACCCTTTGCACAGGTGCTTTCAATAGCTTTTAAATATATTTTTTAAATATATATTT | 60 |
| Qy | 3996 | TAGCTTTTTTAATAAACAAAAATAAATAGCTCTCTTGCTATTTTGGTTTTTGCAAAAG | 4055 |
| Db | 61 | TAGCTTTTTTAATAAACAAAAATAAATAGCTCTCTTGCTATTTTGGTTTTTGCAAAAG | 120 |
| Qy | 4056 | ACCCATCTCAAGGAATGCTGATGTGCTATTAAAAATTTGTTCCAAATGTCCAT | 4109 |
| Db | 121 | ATCCCATCTCAGGAATGCTGATGTGCTATTAAAAATTTGTTCCAAATGTCCAT | 174 |

RESIST 10

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|---------------------------|-------|--------------------|------------|--------------|
| Query Match | 3.2% | Score 165.2; | DB 3; | Length 2873; |
| Best Local Similarity | 60.4% | Pred. No. 6.4e-29; | | |
| Matches 343; Conservative | | 0; Mismatches 193; | Indels 32; | Gaps 3; |

3267 GAGATGCTCAGGTAACCTCAGCAATATGTTGCTCTGCTCTGAAACAACCTAGCGCTGCA 11788
3268 CTTGCTCAGCTGATTTTGTATCCGAAAAAGAAACCCAGGTGATGCTGGGAAGAGAGAGCTG 33273
1179 CCAGGACAGTTTAATATTAAATTTCTAAAGAAAAATAACAAGTGGTGGTGGCAGAGAGAGTTA 12388
Qy Db Qy Db

HOFFMAN, No. 6709821h
KAY, Brian K.
FOWLES, Dana M.
McCONNELL, Stephen J.
TITLE OF INVENTION: POLYPEPTIDES HAVING A FUNCTIONAL
DOMAIN OF INTEREST AND METHODS OF IDENTIFYING AND
USING SAME
NUMBER OF SEQUENCES: 227
CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds LLP
STREET: 1155 Avenue of the Americas
CITY: New York
STATE: New York
COUNTRY: USA
ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/879,957
FILING DATE: 13-Jun-2001
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/630,915
FILING DATE: 03-APR-1996
ATTORNEY/AGENT INFORMATION:
NAME: Misrock, S. Leslie
REGISTRATION/DOCKET NUMBER: 1101-174
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-8864/9741
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 39:
SEQUENCE CHARACTERISTICS:
LENGTH: 747 bases
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: DNA
SEQUENCE DESCRIPTION: SEQ ID NO: 39:
US-09-879-957-39
Query Match 3.2%; Score 163.8; DB 3; Length 747;
Best Local Similarity 63.7%; Pred. No. 6.8e-29;
Matches 277; Conservative 0; Mismatches 137; Indels 21; Gaps 1;
QY 3208 GAAATTGCCAGGTTATGCTCATACACCGCCACCGCCCGCCGAGCAGCTCACTCTCGCC 3267
DB 334 GAGATTGCTCAGGTAACCTCAGCATATGTTGCTTCTGTTCTGACAACTTAGCCTTGCA 393
QY 3268 CCTGGTCAGCTGATTTTGATCCGAAAAAGAACCCAGGTGGATGGTGGGAAGAGAGCTG 3327
DB 394 CCAGGACAGTTAATATTAATTTCTAAAGAAAAATACAAGTGGTGGTGGCAAGAGAGTTA 453
QY 3328 CAAGACGTGGGAAAAAGCCCGCAGATAGGCTGTTCCACGTAATATGTTAAAGCTTCTA 3387
DB 454 CAGGCCAGAGAAAAAGCGACAGAAAGGATGTTCTCCAGTCATGTTAAACTTTTG 513
QY 3388 AGCCCTGGGACGAGCAAAATCACTCCCAACAGAGCCACCTAAGTCAACAGCATTTAGCGCA 3447
DB 514 GGTCCAAGCAGTGAAGAGCCA-----CACCTGCTTTTCATCCT 552
QY 3448 GTGTGCCAGGTGATTTGGATGTACGACTACACCGCCGAGATGACGATGAGCTGGCCTTC 3507
DB 553 GTATGTCCAGGTGATTTGCTATGATGACTATGCGACAAATAATGAAGATGAGCTCAGTTTC 612
QY 3508 AACAGGCCAGATCATCAACGTCCTCAACAGGAGGACCTGACTGGTGGAAAGAGAA 3567
DB 613 TCCAAGGGACAACTCATTAATGTTTGAACAAAGATGATCCTGATGTTGGTGGCAAGAGAG 672
CITY: New York
STATE: New York
COUNTRY: USA
ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/630,915A
FILING DATE: 03-APR-1996
CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: Misrock, S. Leslie
REGISTRATION/DOCKET NUMBER: 1101-174
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 869-8864/9741
TELEFAX: (212) 869-8864/9741
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 39:
SEQUENCE CHARACTERISTICS:
LENGTH: 747 bases
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: DNA
US-08-630-915A-39
Query Match 3.2%; Score 163.8; DB 3; Length 747;
Best Local Similarity 63.7%; Pred. No. 6.8e-29;
Matches 277; Conservative 0; Mismatches 137; Indels 21; Gaps 1;
QY 3208 GAAATTGCCAGGTTATGCTCATACACCGCCACCGCCCGCCGAGCAGCTCACTCTCGCC 3267
DB 334 GAGATTGCTCAGGTAACCTCAGCATATGTTGCTTCTGTTCTGACAACTTAGCCTTGCA 393
QY 3268 CCTGGTCAGCTGATTTTGATCCGAAAAAGAACCCAGGTGGATGGTGGGAAGAGAGCTG 3327
DB 394 CCAGGACAGTTAATATTAATTTCTAAAGAAAAATACAAGTGGTGGTGGCAAGAGAGTTA 453
QY 3328 CAAGACGTGGGAAAAAGCCCGCAGATAGGCTGTTCCACGTAATATGTTAAAGCTTCTA 3387
DB 454 CAGGCCAGAGAAAAAGCGACAGAAAGGATGTTCTCCAGTCATGTTAAACTTTTG 513
QY 3388 AGCCCTGGGACGAGCAAAATCACTCCCAACAGAGCCACCTAAGTCAACAGCATTTAGCGCA 3447
DB 514 GGTCCAAGCAGTGAAGAGCCA-----CACCTGCTTTTCATCCT 552
QY 3448 GTGTGCCAGGTGATTTGGATGTACGACTACACCGCCGAGATGACGATGAGCTGGCCTTC 3507
DB 553 GTATGTCCAGGTGATTTGCTATGATGACTATGCGACAAATAATGAAGATGAGCTCAGTTTC 612
QY 3508 AACAGGCCAGATCATCAACGTCCTCAACAGGAGGACCTGACTGGTGGAAAGAGAA 3567
DB 613 TCCAAGGGACAACTCATTAATGTTTGAACAAAGATGATCCTGATGTTGGTGGCAAGAGAG 672
QY 3568 GTCAATGACAGAGTGGGCTCTCCCATCCAATTTATGTGAAGCTGACACAGCATGGAC 3627
DB 673 ATCAACGGGTGACTGGTCTCTTCTTCAACTACGTTAAGATGACGACAGACTCAGAT 732
QY 3628 CCAAGCCAGCAATGA 3642
DB 733 CCAAGTCAACAGTGA 747
RESULT 13
US-09-879-957-39
Sequence 39, Application US/09879957
Patent No. 6709821
GENERAL INFORMATION:
APPLICANT: SPARKS, Andrew B.

QY 3568 GTCAATGGACAAGTGGGGCTCTTCCCATCCAAATTATGTGAAGCTGACCAACAGACATGGAC 3627
|||||
Db 673 ATCAACGGGGTCACTGGTCTCTTCTTCCAACTAGCTTAAGATGACGACAGACTCAGAT 732
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QY 3628 CCAAGCCAGCAATGA 3642
|||||
Db 733 CCAAGTCAACAGTGA 747
|||||

RESULT 14

US-09-404-879A-5
; Sequence 5, Application US/09404879A

; Patent No. 6486546

; GENERAL INFORMATION:

; APPLICANT: Mitcham, Jennifer L.

; APPLICANT: King, Gordon E.

; APPLICANT: Algate, Paul A.

; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND

; FILE OF INVENTION: DIAGNOSIS OF OVARIAN CANCER

; FILE REFERENCE: 210121.462C2

; CURRENT APPLICATION NUMBER: US/09/404,879A

; CURRENT FILING DATE: 1999-09-24

; NUMBER OF SEQ ID NOS: 393

; SOFTWARE: FastSeq for Windows Version 3.0

; SEQ ID NO 5

; LENGTH: 531

; TYPE: DNA

; ORGANISM: Homo sapien

US-09-404-879A-5

Query Match 3.0%; Score 153.4; DB 3; Length 531;

Best Local Similarity 61.1%; Pred. No. 1.7e-26;

Matches 266; Conservative 0; Mismatches 166; Indels 3; Gaps 1;

QY 175 GCGCTCGATTAGCAAGTAAAGTAACAGAACCATGGCTCAGTTTCCAAACACTTTTGGT 234
|||||
Db 9 GCGTGAAGCTGCAAGAGAGTCAGGATCATGTGGCTCAGTTTCCACAGGGATGAAT 68
|||||
QY 235 GCGAGCTGGATATCTGGGCCATACTGTAGAGGAAGAGCGAAGCATGATCAGCAGTTTC 294
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Db 69 GGAGGCCCAATATGTGGGCTATTACATCTGAAGAACGTACTAAGCATGATAAACAGTTT 128
|||||
QY 295 CATAGTTTAAAGCCAATATCTGGATTCAATCTGATGATCAAGCTAGAACTTTTTTTT 354
|||||
Db 129 GATAACCTCAAACTTTCAGGAGGTTTACATAACAGGTGATCAAGCCCGTACTTTTTCCTA 188
|||||

QY 355 CAATCTGGGTACCTCAACTGTTTTAGCACAGATATGGCCTAGCTGACATGAATAAT 414
|||||
Db 189 CAGTCAGGTCTGCGGCCCGGTTTTAGCTGAAATATGGCCTTATCAGATCTGAACAAG 248
|||||
QY 415 GATGGAAGATGGATCAAGTGGAGTTTTCCATAGCTATGAAACTTATCAAACTGAAGCTA 474
|||||
Db 249 GATGGAAGATGGACCAAGAGAGTTCTCTATAGCTATGAAACTCATCAAGTTAAAGTTG 308
|||||
QY 475 CAAGGATATCAGTACCTCTGCACCTTCCCTCTGTCATGAAACAGCAACC---AGTTGCT 531
|||||
Db 309 CAGGGCCAAACAGCTGCTGTAGTCTCTCTCTCTATCATGAAACAGCCCTTATGTTCTCT 368
|||||

QY 532 ATTTCTAGCGCACCAAGCATTTGGTATGGAGGATATCGCCAGATGCGACCGCTTACAGCT 591
|||||
Db 369 CCACCTAATCTCTGCTCGTTTTGGGATGGAGAGCATGCCCAATCTGTCCATTATCAGCCA 428
|||||
QY 592 GTTGCTCCAGTGCCA 606
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Db 429 TTGCTCCAGTTGCA 443
|||||

RESULT 15

US-09-338-933-5

; Sequence 5, Application US/09338933

; Patent No. 6488931

; GENERAL INFORMATION:

; APPLICANT: Mitcham, Jennifer Lynn

; APPLICANT: King, Gordon E.
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY OF
; FILE OF INVENTION: OVARIAN CANCER
; FILE REFERENCE: 210121.462C1

; CURRENT APPLICATION NUMBER: US/09/338,933

; CURRENT FILING DATE: 1999-06-23

; NUMBER OF SEQ ID NOS: 312

; SOFTWARE: FastSeq for Windows Version 3.0

; SEQ ID NO 5

; LENGTH: 531

; TYPE: DNA

; ORGANISM: Homo sapien

US-09-338-933-5

Query Match

Best Local Similarity 3.0%; Score 153.4; DB 3; Length 531;

Matches 266; Conservative 0; Mismatches 166; Indels 3; Gaps 1;

QY 175 GCGCTCGATTAGCAAGTAAAGTAACAGAACCATGGCTCAGTTTCCAAACACTTTTGGT 234
|||||
Db 9 GCGTGAAGCTGCAAGAGAGTCAGGATCATGTGGCTCAGTTTCCACAGGGATGAAT 68
|||||
QY 235 GCGAGCTGGATATCTGGGCCATACTGTAGAGGAAGAGCGAAGCATGATCAGCAGTTTC 294
|||||
Db 69 GGAGGCCCAATATGTGGGCTATTACATCTGAAGAACGTACTAAGCATGATAAACAGTTT 128
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QY 295 CATAGTTTAAAGCCAATATCTGGATTCAATCTGATGATCAAGCTAGAACTTTTTTTT 354
|||||
Db 129 GATAACCTCAAACTTTCAGGAGGTTTACATAACAGGTGATCAAGCCCGTACTTTTTCCTA 188
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QY 355 CAATCTGGGTACCTCAACTGTTTTAGCACAGATATGGCCTAGCTGACATGAATAAT 414
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Db 189 CAGTCAGGTCTGCGGCCCGGTTTTAGCTGAAATATGGCCTTATCAGATCTGAACAAG 248
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QY 415 GATGGAAGATGGATCAAGTGGAGTTTTCCATAGCTATGAAACTTATCAAACTGAAGCTA 474
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Db 249 GATGGAAGATGGACCAAGAGAGTTCTCTATAGCTATGAAACTCATCAAGTTAAAGTTG 308
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QY 475 CAAGGATATCAGTACCTCTGCACCTTCCCTCTGTCATGAAACAGCAACC---AGTTGCT 531
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Db 309 CAGGGCCAAACAGCTGCTGTAGTCTCTCTCTCTATCATGAAACAGCCCTTATGTTCTCT 368
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QY 532 ATTTCTAGCGCACCAAGCATTTGGTATGGAGGATATCGCCAGATGCGACCGCTTACAGCT 591
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Db 369 CCACCTAATCTCTGCTCGTTTTGGGATGGAGAGCATGCCCAATCTGTCCATTATCAGCCA 428
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QY 592 GTTGCTCCAGTGCCA 606
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Db 429 TTGCTCCAGTTGCA 443
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Search completed: February 14, 2006, 05:36:49

Job time : 845 secs